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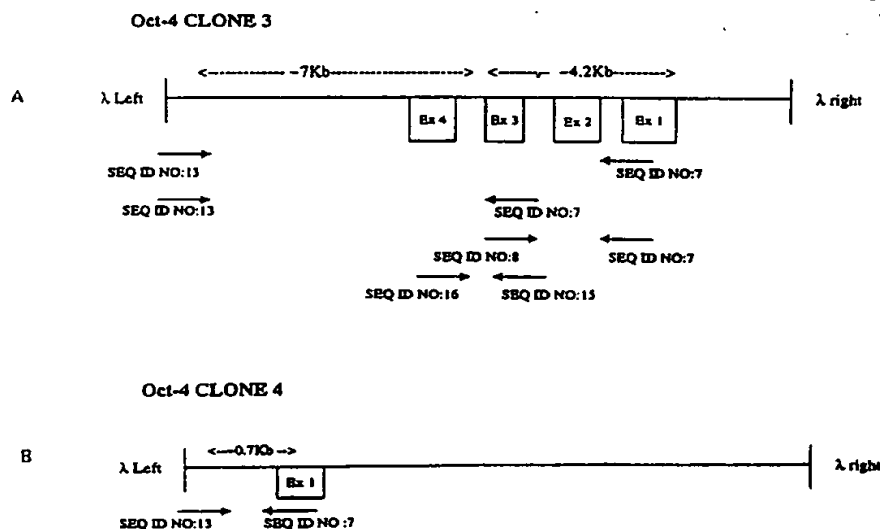
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(54) Title: PORCINE STEM CELLS COMPRISING A MARKER UNDER AN OCT-4 PROMOTER



(57) Abstract

The present invention provides for a method of isolating and/or propagating porcine stem cells, more specifically pluripotent porcine embryonic stem cells. The pluripotent cells are isolated and/or propagated by the use of a selectable marker gene which is inserted into the genetic material of the cells, and which permits the survival and growth of the porcine embryonic stem cells. The selectable marker gene is inserted so as to be regulated by a control or promoter polynucleotide sequence in the embryonic stem cells, for example the promoter polynucleotide sequence being the porcine Oct-4 promoter sequence of the present invention. The invention also provides for a transgenic pig which will constitute a source of the pluripotent cells.

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PORCINE STEM CELLS COMPRISING A MARKER UNDER AN OCT-4 PROMOTER

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Background Of The Invention

15 This invention relates to a method of isolating and/or enriching and/or selectively propagating pluripotential porcine cells, genetically modified porcine cells and pigs for use in said method, transgenic pigs providing a source of such cells and genetic selectable marker constructs for producing genetically modified cells and transgenic pigs.

20 Stem cells are progenitor cells which have the capacity both to self-renew and to differentiate into mature somatic cells. Embryonic stem cells are the archetypal stem cell, being capable of differentiating to form the whole gamut of cell types found in an adult animal. Such stem cells are described as "totipotent" or "pluripotent" as they are capable of differentiating into many cell types. Other types of stem cells, for example bone marrow stem cells and epidermal stem cells, 25 persist in the adult animal. These stem cells have a more restricted capacity for differentiation.

In general, when required for research purposes or for medical use, stem cells have to be isolated from tissue samples by various fractionation procedures. However, even after careful segregation of cell types these stem cell preparations

consist of mixed cell types, and while enriched for stem cells include high proportions of differentiated cells which are not categorized as stem cells.

Furthermore, most stem cells cannot be grown readily in culture. When attempts are made to culture stem cells, the cells being cultured (which ordinarily contain a mixed population of cell types) grow at different rates and stem cells rapidly become overgrown by non-stem cell types. An exception is that embryonic stem cells from two specific strains of mice (129 and Black 6) can be cultured *in vitro* (Evans et al. (1981) Nature 292:154-156). Thus, established lines of murine embryonic stem cells can be obtained by culturing early (3 ½ day) embryonic cells from murine strain 129 and Black 6, or hybrids thereof. Embryonic cell lines from species other than the mouse are not so easily propagated. For an extensive review of isolation and propagation of stem cells see PCT publication WO 94/24274, incorporated by reference herein.

There has developed a pressing need to isolate and maintain *in vitro* embryonic stem cells from species of animals other than murine, such as other laboratory animals and domesticated animals, and most especially, from pigs. However, prior to the present invention the problems associated with producing cultures of porcine (or pig) stem cells, including the problem of producing cell populations of a satisfactorily low degree of heterogeneity and the problem of overgrowth in culture of non-pluripotent porcine cells, have not been solved. A particular problem associated with the continuing presence of certain differentiated cell types is that these can cause elimination of stem cells from the culture by inducing their differentiation or programmed cell death.

Thus according to the present invention, there is provided a porcine cell capable of being cultured under appropriate selective culture conditions so as to enable selective propagation of pluripotential stem cells, characterized in that said pluripotent porcine cells contain a genetic selectable marker, whereby a gene product associated with the genetic selectable marker is produced and which under said culture conditions causes selective survival and/or division of the desired pluripotent cells to occur. "Selective culture conditions" are those conditions under which a

population of cells is selectively grown. For example, to selectively grow cells that contain a gene which transfers resistance to a specific drug, the selective culture conditions would contain the drug so that all cells that did not express the drug resistance would be eliminated. Such selective culture conditions are well known in the art.

The invention further provides according to another aspect thereof, a transgenic animal, in this instance a transgenic pig, having genetic characteristics such that it or its progeny, during embryonic development or later life, constitute a source of porcine pluripotent cells as defined above. Such transgenic pigs may be produced according to the invention by introducing a genetic selectable marker into a fertilized oocyte or an embryonic cell, the genetic marker having the characteristics defined below, and utilizing the resulting transformed oocyte or embryonic cells as a progenitor cell for the desired transgenic animal.

A further aspect of the invention is vectors for use in producing an animal cell, for example a pig cell. Thus the invention further provides vectors for use in genetically modifying animal cells so as to produce transformed cells suitable for use as the source of cells for the method referred to below, said vector comprising a first genetic component corresponding to a genetic selectable marker and a second genetic component which, in the genetically modified porcine cell(s), results in the differential expression of the genetic selectable marker as a stably integrated construct. Such vectors may be in the form of expression vectors in which said second genetic component includes control sequences which are differentially activated in pluripotent stem cells and in cells other than the desired stem cells. The invention covers vectors which when used to transform porcine stem cells become integrated into the animal genome as well as vectors which do not become so integrated.

Summary Of The Invention

The present invention provides for a method of isolating and/or propagating porcine stem cells, more specifically pluripotent porcine embryonic stem cells. The pluripotent (or "pluripotent") cells are isolated and/or propagated by the use

of a selectable marker gene or nucleic acid sequence which is inserted into the genetic material of cells contained in a cell culture comprising porcine pluripotent embryonic stem cells, and which permits the survival and growth of said porcine embryonic stem cells. The selectable marker gene or nucleic acid sequence is
5 inserted so as to be regulated by a control or promoter nucleotide sequence in said embryonic stem cells, for example the control sequence being the porcine Oct-4 promoter nucleotide sequence as described below. The terms "Oct-4 promoter nucleotide sequence" or "Oct-4 promoter sequence" refer to the promoter region of the Oct-4 gene, or any fragment of the promoter region that maintains promoter
10 activity. The invention also provides for a transgenic pig which will constitute a source of said pluripotent cells.

By providing a sufficient and reliable source of porcine pluripotential embryonic stem cells, the present invention permits those skilled in the art to genetically modify the cells with a desired genetic modification. For example, said
15 embryonic stem cells may be genetically altered so as to not express a cell surface membrane protein that may cause rejection of porcine cells after xenotransplantation. Said genetically altered cells are then useful in creating a transgenic pig, or line of transgenic pigs, which will not express said surface membrane protein and which, therefore, will contain organs that are less likely to be rejected upon
20 xenotransplantation.

Brief Description Of The Figures

Figure 1 represents the genomic structure of murine Oct-4 (Yeom et al. (1991) Mechanisms of Development 35:171-179). Figure 1A shows a restriction map of an Oct-4 cosmid derived from the tw5g complete t-haplotype. The position
25 of the transcription unit is indicated with a horizontal arrow running 5' to 3'. The position of the BamHI site missing in wild-type (C3H) and responsible for the restriction fragment length polymorphism (RFLP) between t and wild-type is marked with an asterisk. A vertical arrow indicates a BssHII site; NotI, NruI and MluI sites were absent. Figure 1B shows the exon/intron organization of the Oct-4 transcription
30 unit.

Figures 2A, 2B, and 2C collectively show the polynucleotide sequence of F9 murine Oct-4 cDNA, and the deduced amino acid sequence, of murine Oct-4 protein. The numbering (right for the nucleic acid sequence; left for the amino acid sequence) begins at the putative initiation codon. The POU-specific domain and the POU-homeodomain are marked above the nucleotide sequence.

Figures 3A and 3B collectively show the polynucleotide sequence of a portion of porcine Oct-4 exon 1 determined from genomic porcine DNA obtained from a commercial source (Clontech).

Figures 4A and 4B collectively show the polynucleotide sequence of a contiguous portion of porcine Oct-4 exon 1 determined from genomic porcine DNA from d/d haplotype miniswine.

Figure 5A illustrates the results obtained from PCR mapping of the d/d haplotype miniswine Oct-4 exon 1-containing Lambda clones #3.

Figure 5B illustrates the results obtained from PCR mapping of the d/d haplotype miniswine Oct-4 exon 1-containing Lambda clone #4..

Figures 6A and 6B collectively show the polynucleotide sequence of the porcine Oct-4-related sequence derived from Lambda clone #4.

Figure 7 shows the relationship between two fragments of porcine Oct-4, fragments 5D and 6A, derived from Lambda clone #3, and clone 12 (a PCR product encompassing a portion of exon 1 through exon 3). The letters A through H correspond to oligonucleotides used to map the fragments 5D and 6A derived from Lambda clone #3 and clone 12. Letter A corresponds to SEQ ID NO:7, letters B through H correspond to SEQ ID NO:17 through SEQ ID NO:23. The lengths of fragments derived for the various oligonucleotide pairs are as follows: G→F, 3.5Kb; A→G, 2.2Kb; F→H, 1.2Kb; B→F, 0.9Kb; D→E, 3Kb; H→E, 2.3Kb; and H→C, 3.5Kb.

Figures 8A, 8B, and 8C collectively show the promoter polynucleotide sequence for approximately 3 Kb 5' to the translation initiation codon for porcine Oct-4.

Figures 9A, 9B, and 9C collectively show an alignment comparison of the human, mouse and porcine Oct-4 promoter polynucleotide sequences. The sequence for the human Oct-4 promoter region includes nucleotides 1-499 from Genbank Accession Number Z11900; the sequence for the mouse Oct-4 promoter region includes nucleotides 1401-1950 from Genbank Accession Number S58422S1; the sequence for the porcine Oct-4 promoter region includes nucleotides 2701-3215 from SEQ ID NO:24. The major capping sites (RNA initiation sites) for the murine sequence (Okazawa et al. (1991) EMBO J. 10:2997-3005) are at nucleotides 480 and 501. The box marked as "A" is the SP1/HRE domain.

Figures 10A, 10B, and 10C collectively show an alignment comparison of the Retinoic Acid Responsive Element (RARE) regions of the pig and mouse Oct-4 promoter regions. The mouse sequence also shows high sequence identity within the region corresponding to S58422S1 nucleotides 430 through 1168 and porcine Oct-4 promoter sequence nucleotides 1534 through 2347. This region includes the retinoic acid responsive element located between nucleotides -1132 through -889 of Okazawa et al. (supra) (corresponding to nucleotides 609 through 1101 of Genbank Accession Number S58422S1). This region is also known as the "proximal enhancer region".

Detailed Description Of The Invention

There is an ever growing need for a sufficient and reliable source of pluripotential porcine embryonic stem cells. A particular desire for porcine embryonic stem cells is grounded in the need to be able to genetically modify the stem cells so that later cell development will result in an animal (pig) which will contain organs capable of being accepted immunologically by a recipient host. Once one is able to culture a large number of porcine stem cells and expand them in that state, then they can more easily be genetically modified to create a line of cells which will mature into organs less likely to be rejected after transplantation. These cells

can, according to the invention, be implanted to create a transgenic pig (and a pig line) which will contain organs useful for transplantation into a host.

Donor organ shortages have led to hopes that xenotransplantation could serve as an alternative means of organ availability. Swine, particularly miniswine, are an attractive alternative to nonhuman primate donors because of potentially greater availability, the reduced risk of zoonotic infections, appropriate size of organs and the reduced social and ethical concerns. However, one of the major barriers to xenotransplantation is hyperacute rejection. This phenomenon describes a very rapid and severe humoral rejection, which leads to destruction of the graft within minutes or hours of the transplant of the donor organ. Hyperacute rejection is apparently mediated by a complex series of events, including activation of the complement systems, activation of blood coagulation proteins, activation of endothelial cells and release of inflammatory proteins.

The hyperacute rejection process is initiated when the natural antibodies of the recipient bind to cells of the donor organ. The major cell surface protein (or epitope) that is recognized on porcine cells by human antibodies is the Gal α 1,3Gal β 1,4GlcNAc structure. This structure is expressed at high levels on all mammalian cells, including swine cells, with the exception of human and old world non-human primates. A specific transferase, namely α 1,3 galactosyltransferase, is responsible for the transfer of a terminal galactose to the terminal galactose residue of N-acetyllactosamine-type carbohydrate chains and lactosaminoglycans. If one could eliminate or prevent the expression of the Gal α 1,3Gal β 1,4GlcNAc epitope from the surface of the porcine cells in a transgenic pig, the cells would less likely be rejected by human antibodies and thus organs from such a transgenic pig would more likely be accepted upon transplantation.

In order to eliminate the α 1,3 galactosyltransferase activity in the pig, it would be highly desirable to have an abundant source of porcine embryonic stem cells, in order that sufficient genetic manipulations of the cells could be performed successfully. Using cultured porcine embryonic stem cells, a mutation, preferably a null mutation, is introduced by gene targeting at the native genomic locus encoding

α 1,3 galactosyltransferase. In order to permit such genetic manipulations in a desired quantity, porcine embryonic stem cells must be propagated in large amounts. Once the genetic manipulation is achieved (i.e., the α 1,3 galactosyltransferase activity is eliminated), the stem cells may be utilized in making a line of transgenic pigs. Such transgenic pigs would contain the genetically manipulated cells, which would not express the α 1,3 galactosyltransferase activity, and would therefore provide a supply of organs available for xenotransplantation with a minimized risk of rejection.

The isolation of embryonic stem cells from pigs would provide a method for the multiplication of animals with desired characteristics, an efficient means of producing transgenic animals, and a valuable model for studying cell development and differentiation. One of the major problems associated with the derivation of embryonic stem cells is that most stem cells cannot be grown readily in culture and when attempts are made to culture stem cells, the cells being cultured (usually a heterogeneous cell type population) grow at different rates and stem cells rapidly become overgrown by non-stem cell types. The present invention provides a method for isolating and/or enriching and/or selectively propagating pluripotential porcine embryonic cells which comprises maintaining a source of said cells under culture conditions conducive to cell survival, characterized in that the source of cells includes cells containing a genetic selectable marker which is operatively linked to a porcine promoter nucleotide sequence which provides differential expression of the selectable marker in embryonic stem (pluripotential) cells and cells other than the desired stem cells, whereby differential expression of said genetic selectable marker results in preferential survival and/or division of the desired pluripotential porcine cells. Cell cultures and culture conditions suitable for propagating cells of the present invention are known by those skilled in the art. Suitable cell cultures may be, for example, found in Wurst, W. and A.L. Joyner, "Production of Targeted Embryonic Stem Cell Clones", in Gene Targeting: A Practical Approach (Ed. A.L. Joyner) 1993, Oxford Univ. Press; and Hashimoto et al., WO 95/34636.

In carrying out the method of the invention, the source of cells may include pluripotential cells containing a positive selectable marker and expression of the

marker is used to permit isolation and maintenance of the pluripotent cells. Alternatively (or additionally), the source of cells may include a negative selectable marker which is expressed in cells other than the desired pluripotent cells and is used to deplete the source of cells other than the desired pluripotent cells. The genetic selectable marker may, for example, be a foreign gene, a cellular gene or an antibiotic resistance gene such as for example the bacterial neomycin resistance gene.

It is preferred that the genetic selectable marker is operatively linked to a promoter nucleotide sequence which is differentially active in pluripotent stem cells and non-pluripotent cells. Promoter sequences may be included in the expression construct prior to introduction into the cells. The genetic selectable marker may be introduced into the source of cells by a variety of means known in the art including, but not limited to, injection, transfection, electroporation or by infection with a viral vector.

Further, the source of cells may be produced by transfection extemporaneously, or the source of cells may be derived from a transgenic animal, e.g., the founder transgenic animal or an animal at least one ancestor of which has had the aforementioned genetic marker introduced into its genetic complement. In such transgenic animals it is possible for the genetic selectable marker to pass down the germ line eventually resulting in the production of progeny, from the tissues of which the required source of cells can be derived.

A wide variety of known gene products may be relied upon for selective isolation and propagation of the desired porcine stem cells, including genetic selectable markers which are designed to protect the desired cells from the effects of an inhibiting factor present in the culture medium. In this instance, the inhibiting factor can, for example, be an antibiotic substance which inhibits growth or reproduction of cultured cells not expressing the gene (i.e., cells other than the desired pluripotent stem cells). Alternatively, the genetic selectable marker may selectively permit the growth of pluripotent stem cells using a marker known in the art. In this instance the marker may encode a growth factor, a growth factor

receptor, a transcription factor, or an immortalizing factor. Alternatively, the selectable marker may be a cell surface antigen or other gene product which allows purification or depletion of expressing cells for example by panning or fluorescence-activated cell sorting (FACS). The invention thus enables porcine stem cell
5 populations to be obtained and/or maintained having a satisfactory degree of homology. Examples of all types of selectable markers discussed above are known in the art.

The present invention permits the development of expression constructs which direct specific expression of genetic selectable markers in porcine stem cells
10 and not in differentiated cell types. Having introduced an expression construct by transfection or via the generation of transgenic animals, stem cells present within mixed cell populations can be isolated by culturing in the presence of the selection agent *in vitro*, or by otherwise manipulating the culture conditions by methods well known in the art.

15 One example of a gene which displays a suitably restricted stem cell expression pattern, and therefore should provide a suitable promoter polynucleotide sequence for use in controlling stem cell specific regulatory elements for the expression of a genetic selectable marker in accordance with the invention, is the Oct-4 gene. Octamer binding transcription factor 4 (hereinafter "Oct-4") is a
20 member of the POU family of transcription factors. Oct-4 transcription, controlled by the Oct-4 promoter polynucleotide sequence, is activated between the 4 and 8-cell stage in the developing embryo and it is highly expressed in the expanding blastocyst and then in the pluripotent cells of the egg cylinder. Transcription is down-regulated as the primitive ectoderm differentiates to form mesoderm and by 8.5 days post
25 coitum is restricted to migrating primordial germ cells. High level Oct-4 gene expression is also observed in pluripotent embryo carcinoma and embryonic stem cell lines and is down-regulated when these cells are induced to differentiate. As a result of the inventors' discovery of the sequence of the porcine Oct-4 promoter polynucleotide sequence, it is now possible to link the porcine Oct-4 promoter
30 polynucleotide sequence to a genetic selectable marker in porcine cells and thus isolate and/or propagate a large number of porcine pluripotent stem cells.

As described herein, Applicants' Oct-4 promoter polynucleotide sequence is useful to control a genetic selectable marker sequence to permit propagation of large amounts of porcine embryonic stem cells. The Oct-4 promoter polynucleotide sequence may be used in its entirety, or a portion or fragment of the promoter sequence may be used in which the portion maintains the promoter activity. One skilled in the art would easily be capable of, using Applicants' sequence, joining a selectable marker to a portion or fragment of the porcine Oct-4 promoter polynucleotide sequence, transfecting the Oct-4-marker construct into a colony of cells, and growing the transfected cells in the appropriate medium to determine if the genetic selectable marker is translated. Within the scope of Applicants' invention are such portions or fragments that retain promoter activity, such portions having preferably at least 90% sequence identity, more preferably 95% sequence identity, and most preferably 98% sequence identity to the porcine Oct-4 promoter polynucleotide sequence as shown in Figure 8 or a portion of said Oct-4 promoter polynucleotide sequence, wherein said portion preferably consists of contiguous nucleotides from the porcine Oct-4 promoter polynucleotide sequence, i.e., a contiguous portion, and wherein said portion preferably comprises at least 38 nucleotides, more preferably at least 100 nucleotides, more preferably at least 200 nucleotides, more preferably at least 500 nucleotides, more preferably at least 1000 nucleotides, more preferably at least 1500 nucleotides, more preferably at least 2000 nucleotides, and most preferably at least 2500 nucleotides of the porcine Oct-4 promoter polynucleotide sequence shown in Figure 8.

Also within the scope of Applicants invention are polynucleotide sequences which hybridize to all or a portion of the Oct-4 promoter polynucleotide sequence as shown in Figure 8. Preferred under the scope of Applicants' invention are polynucleotide sequences which hybridize under high stringency conditions to all or a portion of the porcine Oct-4 promoter polynucleotide sequence as shown in Figure 8. Conditions under which hybridization will occur are known in the art and can be found in, for example, Bulletin 1234, Bio-Rad Laboratories, incorporated by reference herein.

The polynucleotides of the present invention may be in the form of DNA which DNA includes genomic DNA and synthetic DNA. The DNA may be double-stranded or single-stranded, and if single stranded may be either of the strands which together comprise the promoter. Such sequences are useful either in the promoter
5 functionality or as probes to retrieve the promoter sequences.

Fragments of the full-length promoter of the present invention may be used as hybridization probes for the DNA containing the promoter or to isolate other DNAs which have a high polynucleotide sequence identity to the promoter. Probes of this type preferably have at least 10 nucleotides, preferably at least 15 nucleotides, and
10 even more preferably at least 30 nucleotides and may contain, for example, at least 50 or more nucleotides. In fact, probes of this type having at least up to 150 nucleotides or greater may be utilized. An example of a screen comprises isolating the promoter region of the porcine Oct-4 gene by using the known DNA sequence to synthesize an oligonucleotide probe. Labeled oligonucleotides, having a sequence
15 complementary to that of the promoter or portion of the promoter sequences of the present invention are used to identify those polynucleotides that hybridize to, in a complementary sense, the promoter fragment, and have an identity as described above.

It is also appreciated that such probes can be and are preferably labeled with
20 an analytically detectable reagent to facilitate identification of the probe. Useful reagents include but are not limited to radioactive labels, fluorescent dyes or enzymes capable of catalyzing the formation of a detectable product. The probes are thus useful to isolate complementary copies of the Oct-4 related promoter sequences from other sources or to screen such sources for related sequences.

25 The present invention further relates to polynucleotides which hybridize to the hereinabove-described sequences (SEQ ID NO:24 or a polynucleotide sequence encoding the same promoter as encoded by the sequence according to SEQ ID NO:24) if there is at least 70%, preferably at least 90%, and more preferably at least 95% identity between the sequences.

The polynucleotides of the present invention are preferably provided in an isolated form, and preferably are purified to homogeneity.

The term "isolated" means that the material is removed from its original environment (e.g., the natural environment

5 Moreover, within the scope of Applicants' invention is the use of the porcine Oct-4 promoter polynucleotide sequence, or a portion thereof, connected to all or a portion of exon 1 and intron 1 of the porcine Oct-4 gene. The Oct-4 promoter plus exon/intron 1 is then joined to a genetic selectable marker according to the invention to be used in selectively propagating porcine stem cells.

10 Using applicants' teaching, pluripotential porcine embryonic stem cells are created in which the porcine Oct-4 promoter polynucleotide sequence is employed to drive stem cell specific transcription of a selectable marker. The stem cells are propagated in large quantities and are available for further genetic manipulation (e.g., to eliminate the α 1,3 galactosyltransferase activity). Applicants' invention also
15 provides for transgenic pigs in which a porcine Oct-4 promoter polynucleotide sequence drives specific transcription of the selectable marker. An appropriate genetic selectable marker, for example, is the neomycin phosphotransferase gene which confers resistance to the antibiotic G418, although other genetic selectable markers are known and are available to those skilled in the art.

20 Selectable marker genes under the control of the porcine Oct-4 promoter may, according to the invention, be applied to the isolation of embryonic stem cell lineages from a transgenic pig.

 Accordingly, this invention includes the generation of transgenic pigs which express the porcine Oct-4 promoter polynucleotide sequence linked to a genetic
25 selectable marker, for example the selectable marker being β -geo (sequences encoding a fusion protein of β -galactosidase and neomycin phosphotransferase II). The β -galactosidase activity enables visualization of cells expressing the marker and the neomycin phosphotransferase activity confers resistance to the antibiotic G418.

Embryonic stem cells are then isolated from the transgenic pigs using culture conditions to support the growth of undifferentiated cells and maintaining selective pressure for cell expressing β -geo under the control of the Oct-4 promoter polynucleotide sequence.

5 Also included within the scope of the invention are transgenic pigs which contain cells expressing a genetic selectable marker under the control of the porcine Oct-4 promoter polynucleotide sequence according to the present invention, as well as transgenic pigs which contain cells expressing a genetic selectable marker under the control of the porcine Oct-4 promoter polynucleotide sequence which have been
10 genetically manipulated to eliminate the α 1,3 galactosyltransferase activity. The pigs containing the genetically manipulated cells are good resources for organs suitable for transplantation because of the minimal risk of hyperacute rejection.

In order to obtain the porcine Oct-4 promoter polynucleotide sequence, Applicants isolated porcine Oct-4 gene sequences and used those sequences to isolate
15 and determine the DNA sequence upstream of the start of transcription initiation. The genomic map of the mouse Oct-4 gene (Yeom et al. (1991) Mechanisms of Development 35:171-179) is presented in Figures 1A and 1B. The entire sequence of the mouse Oct-4 gene (Schöler et al. (1990) Nature 344:435-439) is contained on a 4.73 Kb BamHI fragment (Figures 2A, 2B, and 2C). There are five exons and four
20 introns. The sizes of the two short introns, 2 and 4, are 182 and 139 bp, respectively; the sizes of introns 1 and 3 are approximately 2430 and 450 bp, respectively. The POU domain is encoded by two or three split exons. The cap site of the murine Oct-4 transcript is located 163 bp upstream of the start codon (Yeom, 1991).

The invention will now be described in more detail in the following non-
25 limiting examples with reference to the drawings. The examples are for illustration only and do not limit the scope of the present invention in any way, which is defined only by the claims. All references cited are hereby incorporated by reference herein in their entirety.

EXAMPLE 1 Cloning of Porcine Oct-4

1.1 Obtaining porcine Oct-4 exon 4 sequences

In order to obtain a fragment of porcine genomic DNA containing the Oct-4 sequence, degenerate PCR primers were designed based on sequence identity to several Oct family genes found in the Entrez database (http://www.ncbi.nlm.nih.gov/). These primers were designed such that they might amplify different Oct family members and were not specific for Oct-4. The sequences of the two degenerate primers are as follows:

DA34 (sense): 5' GCCCCTSCTGGAGAAGTGG 3' (SEQ ID NO:1)

10 DA37 (antisense): 5' GSCGSCGGTTRCAGAACCA 3' (SEQ ID NO:2)

DA34 maps within exon 3, which contains part of the POU-specific domain. DA37 maps within exon 4, which contains the POU-homeodomain.

Using a commercial source of porcine genomic DNA (Clontech, Palo Alto, CA, catalog # 6651-2) as template, a PCR fragment of approximately 1 Kb was amplified using Touchdown PCR (Roux, 1994) with Vent DNA Polymerase (New England Biolabs, Inc. ("NEB") Beverly, MA) on a Perkin-Elmer 9600 Thermocycler (Perkin-Elmer Corp., Norwalk, CT). Following NEB's protocol for setting up PCR reactions, the samples were amplified as follows: The tubes were heated at 94°C for 2 minutes followed by 3 cycles, each consisting of 45 seconds at 94°C, 1 minute at 55°C and 2 minutes at 72°C. The 3-cycle PCR step was repeated for 14 cycles during which the annealing temperature was sequentially decreased by 1°C every cycle. The final annealing temperature after 14 cycles was 40°C. Following this, 10 cycles were carried out at 94°C for 45 seconds, 1 minute at 40°C and 2 minutes at 72°C. A final extension for 5 minutes at 72°C completed the PCR program.

25 The 1 Kb PCR fragment was purified by gel electrophoresis followed by clean up on Qiaex resin (Qiagen, Chatsworth, CA) according to its manufacturer's protocol. The PCR fragment was treated with Taq polymerase in order to add the 3' A overhang. This allowed the cloning of this fragment into the TA vector, pCR II (Invitrogen, San Diego, CA). Several clones were sequenced using the primer

walking technique and standard DNA sequencing methodology. The complete sequence for clone pDA118-1 (960 bp) was determined. Based on sequence identity to both the human and mouse Oct-4, this clone was concluded to be authentic porcine Oct-4. The region of the Oct-4 gene contained within this clone extends from the last 48 bp of exon 3 to the first 27 bp of exon 5. Intron 3 was 628 bp, exon 4 was 159 bp and intron 4 was 99 bp in length.

1.2 Amplification, cloning and partial sequencing of a fragment from porcine genomic DNA which extends from exon 1 to exon 4.

10 A degenerate sense PCR primer within exon 1 was designed based on the sequence of human (Takeda et al. (1992) Nucleic Acids Research 20:4613-4620) and mouse Oct-4 (Okazawa et al. (1991) EMBO J. 10:2997-3005). The primer location was selected within the most conserved region of exon 1:

DA44 (sense): 5' CACCTGGCTTCRGAYTTTCGCCTTC 3' (SEQ ID NO:3)

15

Using the DNA sequence derived from the 960 bp PCR fragment (clone PDA118-1), isolated from porcine genomic DNA (Clontech, Palo Alto, CA) and sequenced as described in 1.1 above, 2 antisense PCR primers were designed which map within either exon 3 or 4, respectively, and contain the following sequences:

20 DA43 (antisense, exon 3): 5' GCAGATTCTCGTTGTTGTCAGCTT 3' (SEQ ID NO:4)

DA58 (antisense, exon 4): 5' GTTGCCTCTCACTCGGTTCTCGATAC 3' (SEQ ID NO:5)

25 Using porcine genomic DNA (Clontech, Palo Alto, CA) as template, 2 PCR reactions were set up with either primer pair DA44/43 (SEQ ID NO:3/SEQ ID NO:4) or DA44/58 (SEQ ID NO:3/ SEQ ID NO:5) using the TaKaRa LA Taq Kit (Pan Vera, Madison, WI). PCR conditions were as recommended by the manufacturer, namely: 94°C for 2 minutes, 30 cycles at 98°C for 15 seconds, then 65°C for 11
30 minutes, followed by a 72°C extension for 5 minutes. The expected sizes of the PCR

products ranged from 3.2 to 6.8 kb. The observed PCR products were ~4.2 Kb for exon 1/3 primers and ~5.1 Kb for exon 1/4 primers.

The PCR products were purified by gel electrophoresis followed by clean-up on Qiaex resin (Qiagen, Chatsworth, CA). The purified fragments were cloned directly into the TA vector pCR II, transformed and bacteria colonies screened. Two clones were selected which contained either exon 1/3 fragment (pDA131-5) or exon 1/4 fragment (pDA132-9). Large-scale DNA preparations were made of each of these clones using Qiagen DNA Kit. Using standard DNA sequencing methodology that included the use of the SequiTherm Cycle Sequencing Kit (Epicentre Technologies, Madison, WI) and their vector sequencing primers, both clones were sequenced in from both ends. Sequence comparison of pDA132-9 with the sequence from pDA118-1 derived in Example 1.1 confirmed that this clone was authentic Oct-4. Exon 3 sequence of the two clones was also very similar. A consensus exon 1 sequence was obtained from sequence data of both clones. Approximately 340 bp of exon 1 sequence was determined for porcine Oct-4, as shown in Figures 3A and 3B (SEQ ID NO:6). This sequence showed a high degree of sequence identity to both human and mouse Oct-4 sequences.

1.3 Obtaining an Oct-4 exon 1 probe

A second porcine Oct-4 probe was generated using the following oligonucleotide primers:

Oct-4 exon 1 5': 5' GGATCCTCGGACCTGGCTGAGCTTCCAA 3' (SEQ ID NO:7)

Oct-4 exon 3 3': 5' GAGCTCGTTGTTGTCAGCTTCCTCCACCCA 3' (inverse complement) (SEQ ID NO:8)

The oligonucleotide primer SEQ ID NO:7 includes nucleotides 92 through 118 of SEQ ID NO:6, while SEQ ID NO:8 was derived from the porcine exon 3 sequence.

Since the size in both the human and murine counterparts of porcine Oct-4, intron 1, is known to be quite large, i.e., 5 Kb and 2.43 Kb respectively (Takeda,

1992; Yeom, 1991), the PCRs were carried out using the TaKaRa LA Taq kit (Pan Vera Corp., Madison, WI). Each 50µl reaction contained 5 µl 10X LA PCR buffer II, 5 µl 25 mM MgCl₂, 8 µl dNTP mix (2.5 mM each), 2.5 units of TaKaRa LA Taq (0.5 µl), 500ng template DNA (porcine genomic DNA from Clontech (Palo Alto, CA) or from d/d miniswine #s 12021, 11378, 12023, 12037, 12038) and 100 pmol each oligonucleotide. The polymerase chain reaction (PCR) was performed using a Perkin Elmer DNA Thermal Cycler 480. The program used included 5 minutes at 94°C, followed by 14 cycles, each containing 20 seconds at 98°C and 8 minutes at 68°C. This was followed by 16 cycles, each consisting of 20 seconds at 98°C and 8 minutes at 68°C with an autoextension time of 15 seconds/cycle. The program concluded with a final extension of 5 minutes at 72°C and a 4°C soak. All reactions, except one, produced two PCR products: a major band at approximately 4.5 Kb and a minor band at 0.6 Kb (miniswine # 12037 gave only the smaller band). These were the sizes one would expect from an intron containing gene and an intronless gene, respectively, within this portion of Oct-4.

The 4.5 Kb PCR fragment obtained from miniswine #12038 was gel isolated using Qiaex II resin (Qiagen Inc., Chatsworth, CA) cloned into the vector pCR II and the resultant plasmid transformed into *E. coli* Inv α F' using the instructions provided with the TA cloning kit (Invitrogen, San Diego CA). In order to identify those clones which contained the Oct-4 insert, two oligonucleotides were ordered from Genosys Biotechnologies (The Woodlands, TX) and colony PCRs were performed. The two oligonucleotides are shown below:

Oct-4 Ex 1- 5' seq: 5' CGGACCTGGCTGAGCTTCCAA 3' (SEQ ID NO:9)

Oct-4 Ex 1- 3' seq: 5' CCTCGGAGTTGCTCTCCACC 3' (SEQ ID NO:10)

SEQ ID NO:9 contains nucleotides 98 through 118 of SEQ ID NO:6, and SEQ ID NO:10 contains the reverse complement of nucleotides 307 through 326 of SEQ ID NO:6.

Bacteria derived from each of 40 white (presumed to be insert containing) colonies were separately inoculated into 50 µl of distilled water. The 40 samples were boiled for 5 minutes and centrifuged briefly in order to pellet the cells. Ten µl of each supernatant then served as template for PCRs which were carried out as

described previously for Taq DNA polymerase with the following exceptions: 1) the samples contained 50 µl rather than 100 µl and 2) the program used was 5' at 94°C, followed by 35 cycles, each containing 1 minute at 94°C, 1 minute at 55°C and 1 minute at 72°C. There was a final extension of 10 minutes at 72°C followed by a 4°C soak. The two oligonucleotides used in the PCRs should amplify a 230 bp exon 1 fragment of Oct-4. Most of the colonies tested were positive and one (#12) was chosen for sequence analysis. All sequencing was performed using either the Sequenase version 2.0 kit (United States Biochemical Corp., Cleveland, Ohio 44122) or the Fidelity DNA Sequencing kit (Oncor, Inc. Gaithersburg, MD 20877). Figures 4A and 4B (SEQ ID NO:11) show the sequence of the 5' end of clone #12, which includes most of exon 1 and a small portion of intron 1. By sequence comparison to the human and murine sequences, the exon/intron junction is estimated to be at nucleotide 320. The amino acid sequence of exon 1, which shows strong homology to both the murine and human genes, is underlined.

In order to prepare an exon 1 probe for use in a genomic Southern as well as in a subsequent library screen, the oligonucleotide shown above as SEQ ID NO:10 and the following oligonucleotide (SEQ ID NO:12) were used in a PCR for which the 4.5 Kb Oct-4 insert of clone #12 served as template:

Oct-4 ex 1 probe 5': 5' GGATCCTCGGACCTGGCTGAGC 3' (SEQ ID NO:12)

SEQ ID NO:12 includes nucleotides 92 through 112 of SEQ ID NO:6.

Ten identical PCRs, using Taq DNA polymerase, were carried out as described previously. A small portion of each reaction product was run on a 1% agarose gel in order to confirm the presence of a 236 bp band. The remainder of the reactions were then pooled, chloroform extracted, and ethanol precipitated. After resuspension in TE (10 mM Tris-HCl, pH 8.0, 1 mM EDTA), the DNA fragment was quantitated by running a portion on a 1% agarose gel along with markers containing known quantities of DNA.

1.4 Genomic Southern to verify utility of Oct-4 exon 1 probe

A genomic Southern was performed in order to determine whether or not the 236 bp probe, described in the previous section, would hybridize with specificity and thus could be safely used in a library screen (see Molecular Cloning, a Laboratory Manual, second edition, J. Sambrook et. al., Cold Spring Harbor Laboratory Press, 1989). The genomic DNA used in both the Southern blot and library construction was obtained from d/d miniswine # 11852. For the Southern, 20 µg/reaction of genomic DNA was digested at 37°C overnight in a 50 µl volume with each of the following restriction enzymes: Bam HI, Bgl II, Eco RI, Hind III, Pst I, Sac I, Xba I (New England Biolabs, Inc., Beverly, MA). A small sample of each digest was run on a 0.7 % agarose gel in order to insure that the digests were complete. The remainder of each of the 7 reactions was then ethanol precipitated, rinsed in 70 % ethanol, dried, resuspended in 25 µl TE and run on a 15 X 15 cm 0.7 % agarose gel overnight at 20 V. A positive control was also included on 2 lanes of the gel, i.e., 2 pg and 20 pg respectively of the 4.5 Kb Oct-4 insert obtained by PCR, as previously described.

The next morning, the gel was soaked in 0.25 N HCl for 10 minutes, rinsed in water, and the DNA denatured for 45 minutes by soaking the gel in 1.5 M NaCl, 0.5 M NaOH. After a water rinse, the gel was placed for 45 minutes in a neutralization solution containing 1.5 M NaCl, 1M Tris-HCl pH 7.4 and then for another 45 minutes in 10X SSC (1.5 M sodium chloride/0.15 M sodium citrate, pH 7.0). The DNA was transferred onto a nitrocellulose membrane in 10X SSC overnight using the TurboBlotter Rapid Downward Transfer System (Schleicher & Schuell, Inc., Keene, NH 03431). After transfer, the nitrocellulose membrane was rinsed for 5 minutes in 5X SSPE (0.75 M sodium chloride, 0.05 M sodium phosphate, 0.005M EDTA) and then baked for 2 hours at 80°C in a vacuum oven.

Approximately 50 ng of the 236 bp probe was labeled by random priming with α -³²P dCTP according to instructions provided with the High Prime DNA Labeling Kit (Boehringer Mannheim, Indianapolis, IN). The nitrocellulose blot was prehybridized at 65°C for 3 hours in a solution containing 6X SSC, 5X Denhardt's (0.5 g Ficoll, 0.5 g polyvinylpyrrolidone, 0.5 g bovine serum albumin), 100 µg/ml

sheared denatured herring sperm DNA, 0.75 % sodium dodecyl sulfate (SDS).

Hybridization was carried out overnight at 65°C in the same solution to which the probe was added, following its denaturation in 1/10 volume 0.1 N NaOH. The blot was then washed 2 X 15 minutes at room temperature in 7X SSPE, 0.5 % SDS, 2 X 15 minutes at 37°C in 1X SSPE, 0.5 % SDS, 1 X 15 minutes at 65°C in 0.1X SSPE, 1 % SDS, rinsed in 0.1X SSPE at room temperature and exposed to X-ray film for 2 days at -80°C.

The results indicated that the Oct-4 exon 1 probe was quite specific since only 1 or 2 bands appeared for each of the restriction digests performed.

Hybridization of the probe to the positive control was also seen, but only in the lane which contained the higher concentration (20 pg) of DNA.

1.5 Construction of a miniswine haplotype d/d genomic library

Miniswine genomic DNA (11852, haplotype d/d) was extracted from liver tissue and partially digested with Sau3AI. The Sau3AI sites were then partially filled-in and the fragments were size fractionated by gel electrophoresis using standard methodology. The genomic fragments were ligated to LambdaGEM-12 XhoI partially filled-in according to manufacturer's instructions (Promega, Madison, WI), packaged using Stratagene's Gold extract and titered on E. coli KW251 host cells.

The genomic library had 1.5×10^6 independent clones, with an average insert size of ~12 Kb. The library was amplified one time, aliquoted and stored. The final titer of the amplified library was 1×10^{10} pfu/ml.

1.6 Screening the miniswine haplotype d/d genomic library for Oct-4 exon 1-containing clones

Approximately 3×10^5 clones were screened with the Oct-4 exon 1 probe using duplicate nitrocellulose filters (Sambrook, 1989). The prehybridization, hybridization and wash conditions were essentially the same as those used for the Southern blot. The first round screen produced 5 positives; each of these was

replated at 3 different plaque densities and screened again with the Oct-4 exon 1 probe. Only 2 of the 5 were still positive after the second round screen. The two remaining clones (designated as #3 and #4 respectively) were plaque purified after a third round screen and λ DNA was prepared from plate lysates using a Qiagen
5 Lambda Kit (Qiagen, Chatsworth, CA).

1.7 Mapping the Oct-4 exon 1-containing clones by PCR

Lambda clones #3 and #4 were mapped using LA PCR (Pan Vera, Madison, WI). The reactions were carried out as described previously except for a change in
10 the program which increased all times at 68°C from 8 to 12 minutes. Two new oligonucleotides were made based on sequence provided by Promega (Madison, WI), the supplier of the LambdaGEM-12 vector:

RK EMBL3L: 5' GCAACGAACAGGTCACTATCAGTCA 3' (left arm of vector) (SEQ ID NO:13)

15 RK EMBL3R: 5' CTGCCTTCATTAAGGGCTGCGCAC 3' (right arm of vector) (SEQ ID NO:14)

It should be pointed out that the oligonucleotide shown as SEQ ID NO:14 never gave a positive PCR result, nor did two other oligonucleotides designed to
20 amplify portions of the Oct-4 clones from the right arm of the vector. As can be seen in Figure 5, both clones are inserted in the vector with their 5' ends toward the right arm and their 3' ends toward the left arm of LambdaGEM-12. When the oligonucleotide pair of SEQ ID NO:7 and SEQ ID NO:13 was used, the size of the fragment amplified from clone #3 was approximately 11 Kb while the same pair
25 amplified a much smaller 0.7 Kb fragment from clone #4. Additional PCRs using oligonucleotide pairs SEQ ID NO:15 and SEQ ID NO:13, SEQ ID NO:7 and SEQ ID NO:8, or SEQ ID NO:15 and SEQ ID NO:16 generated 7 Kb, 4.2 Kb, and 0.9 Kb products respectively from clone #3, but no products at all from clone #4. SEQ ID
30 NO:15 includes sequences present in the porcine Oct-4 exon 4. SEQ ID NO:16 includes sequences in the exon 5/intron 4 junction region of porcine Oct-4.

Oct4 exon 3 5': 5'CTCGAGAAAGTGGGTGGAGGAAGCGAC 3' (SEQ ID NO:15)

Oct4 exon 5/intron 4 3' : 5'GAATTCCACGCGGACCACCTGAGAG
CAGGA3' (reverse complement) (SEQ ID NO:16)

These PCR results indicated that clone #3 contained the entire coding region of Oct-4 (i.e. exon 1 through exon 5 in addition to all 4 introns), several kilobases downstream, plus 2-3 kilobases upstream; clone #4 appeared to consist mostly of exon 1 and upstream regions.

1.8 DNA Sequence Analysis of Lambda Clones #3 and #4

In order to obtain exon 1 containing fragments of Lambda clones 3 and 4 for DNA sequence analysis, a Southern blot was prepared and probed with the exon 1 probe described earlier. The following restriction enzymes were used to digest 1-2 µg of clones 3 and 4 respectively, in 25 µl reactions: Bam HI, Eco RI, Hind III, Sac I, Xba I. All digests were carried out for 6 hours at 37°C and then loaded directly onto a 15 X 15 cm 1% agarose gel. The gel was run overnight at 20 V and the DNA transferred as described previously with the following changes: 1) instead of nitrocellulose, a positively charged nylon membrane was used, and 2) transfer was carried out in 20 X SSC instead of 10 X SSC.

The 236 bp exon 1 probe was labeled non-radioactively by random priming with DIG-11-dUTP. Labeling was carried out according to instructions provided with the Genius-2 DNA Labeling Kit (Boehringer Mannheim Corp., Indianapolis, IN). Following transfer, the DNA was affixed to the nylon blot by irradiation in a U.V. 1800 Stratalinker (Stratagene, La Jolla, CA). The blot was prehybridized for 3 hours at 50°C in DIG Easy Hyb (Boehringer Mannheim) and then hybridized overnight at 50°C in 15 ml of the same solution to which approximately 115 ng of boiled, denatured DIG-labeled probe had been added. Following hybridization, the blot was washed 2 X 15 minutes at room temperature in 2X SSC, 0.1% SDS, and then 2 X 15 minutes in 0.1X SSC, 0.1% SDS at 50°C. Oct-4 exon 1 containing bands were detected by chemiluminescence using the reagents Anti-Digoxigenin-alkaline phosphatase and the substrate CSPD, following instructions obtained from
Boehringer Mannheim.

Four fragments which hybridized with the exon 1 probe, 2 from each Lambda clone (#3 and #4), were gel isolated using Qiaex resin and ligated into pBluescript KSII for sequencing. The resultant clones are described below:

1. 5D is a 5.5 Kb Bam HI fragment derived from Lambda clone #3. It contains
5 exon 1, a portion of intron 1 and approximately 3 Kb upstream of exon 1.
2. 6A is a 4 Kb Sac I fragment derived from Lambda clone #3 which overlaps
clone 5D. It contains exon 1, slightly more than 1 Kb upstream, and contains 1 Kb
more of intron 1 downstream than does 5D.
3. 9-5 is a 1.3 Kb Hind III fragment derived from Lambda clone #4.
- 10 4. 10-C is a 0.6 Kb Sac I fragment of Lambda clone #4 which overlaps the Hind
III clone above.

Sequence analysis of the overlapping fragments derived from Lambda clone
#4 indicated that they were not derived from authentic Oct-4. The exon 1 sequence
15 was related, but not identical to sequence obtained previously. There was no open
reading frame in this exon 1-like DNA stretch. Furthermore it was joined to a
sequence which was similar (but not identical) to that of exon 2, but with no intron 1
between them. A portion of this sequence can be seen in Figures 6A and 6B. Since
this clone did not appear to contain authentic Oct-4, no further sequence analysis was
20 carried out on λ clone #4.

Figure 7 shows the relationship between fragments 5D and 6A derived from
Lambda clone #3 and clone 12, described earlier. Mapping was done by sequence
analysis as well as by LA PCR, using a number of oligonucleotides (SEQ ID NO:7
and SEQ ID NO:17 through SEQ ID NO:23) which are identified by the letters A
25 through H on Figure 7. As can be seen, the 3 clones represent overlapping segments
of porcine Oct-4 which extend from approximately 3 Kb upstream of exon 1 through
exon 3. DNA sequence obtained from fragments 5D and 6A upstream of exon 1
further confirm that they are derived from authentic Oct-4. The murine counterpart
of porcine Oct-4 (Oct3/4) lacks a TATA box, but possesses a putative Sp1/HRE
30 (hormone-responsive element) in the proximal promoter region (Minucci et. al.
(1996) EMBO J. 15:888-899). An identical sequence (see below) is found
approximately 100 bp upstream of the translational start in the porcine gene, which

also lacks a TATA box: 5' GGGGGCGGGGCCAGAGGTCAAGGCTA 3'. The sequence determined for the promoter polynucleotide sequence of the porcine Oct-4 gene is shown in Figures 8A, 8B, and 8C (SEQ ID NO:24). This is what is referred to herein as Applicants' "Oct-4 promoter polynucleotide sequence" or the "Oct-4 promoter sequence". Figures 9A-C collectively show an alignment comparison of the human, mouse, and porcine Oct-4 promoter oligonucleotide sequences. Figures 10A-C collectively show an alignment comparison of the Retinoic Acid Responsive Element (RARA) regions of the porcine and murine Oct-4 promoter regions.

10 EXAMPLE 2 Testing of the Oct-4 Promoter

In order to demonstrate that the porcine Oct-4 promoter polynucleotide sequence is an embryonic stage specific promoter, the following constructs were made and tested in mouse embryonic stem cells.

2.1 Construction of pOct-4-βGeo

15 Plasmid pGT1.8 was obtained from Dr. P.S. Mountford (Stem Cell Sciences, Melbourne, Victoria, Australia). pGT1.8 contains a nuclear targeted Internal Ribosomal Entry Site (IRES) -βGeo construct. βGeo is a gene fusion of sequences encoding β-galactosidase and neomycin resistance gene (Friedrich et al. (1991) Genes & Development 5:1513-1523). 5' to the IRES is a sequence containing
20 murine engrailed-2 splice acceptor which is designated as EN-2 (Gossler et al. (1989) Science 244:463-465). A 5.5 Kb fragment containing the porcine Oct-4 promoter, exon 1 and a portion of intron 1 was inserted into the SalI site of pGT1.8 as follows: Clone 5D (Example 1.8 above) was incubated with SalI/NotI and the 5.5 Kb DNA fragment was isolated. The DNA fragment was cleaved using Eco47III and the large
25 4.5 Kb 5'SalI/Eco47III DNA fragment was gel isolated.

A 0.6 Kb 3' portion of 5D was isolated after digestion with Eco47III/BamHI. The 3'BamHI site at the end of this fragment was converted to a XhoI site by PCR mutagenesis, using the following oligonucleotides:

PM Eco47III 5" 5' TCAAAGCGCTAAAATGTGATTTGG (SEQ ID NO:25)

30 PM XhoI3' 5' CGATCTCGAGGGATCCCAGACCGGGGA ACT (SEQ ID NO:26)

The resulting PCR product was cloned into the TA vector, pCRII (Invitrogen, San Diego, CA) and sequenced. A correct clone was identified, cleaved with Eco47III and XhoI and gel purified. The final construct was constructed by ligation of dephosphorylated pGT1.8/SalI, the 4.5 Kb SalI/Eco47III 5' portion of 5D, and the
5 0.6 Kb Eco47III/XhoI 3' fragment. Since SalI and XhoI generate compatible overhanging ends, the Oct-4 insert could be inserted into pGT1.8 in either of two orientations. These orientations can be distinguished by XbaI cleavage. There are two XbaI sites in pGT1.8 and three in the Oct-4 insert: 125, 950, and 1990 bp from the SalI site. The correct orientation results in the following size fragments upon
10 XbaI digestion: 5.3 Kb, 4.9 Kb, 3.0 Kb, 1.0 Kb, and 0.8 Kb. XbaI cleavage of the incorrect orientation generates 6.3 Kb, 4.9 Kb, 2.0 Kb, 1.0 Kb, and 0.8 Kb fragments.

2.2 Construction of pPGK- β Geo

The mouse PGK promoter was removed from a PGKneo vector (Aron Thall, BioTransplant, Inc.) using EcoRI and PstI. The resulting 0.5 Kb fragment was gel
15 purified and the overhanging ends were filled in with Klenow DNA polymerase (NEB, Beverly, MA) according to instructions, phenol-chloroform extracted, and ethanol-precipitated. The DNA was resuspended in 1.5 μ l H₂O, and 2 μ g of phosphorylated HindIII linkers (2 μ l) was added together with 1 μ l 10 mM ATP, 1 μ l
20 100 mM DTT, 2 μ l 10X ligation buffer (Novagen, Madison, WI), and 1.5 μ l T4 DNA ligase. The ligation reaction was incubated overnight at 16°C. The resulting DNA was digested with EcoRI and HindIII and gel-purified. pOCUS-2 (Novagen, Madison, WI) was digested with EcoRI and HindIII and desphosphorylated using shrimp alkaline phosphatase (SAP). A ligation was performed with the pOCUS-2
25 fragment and the mouse PGK-promoter fragment. Following transformation of E. coli, colony PCRs were performed using primers that correspond to sequences located on either side of the multiple cloning site (MCS). The PCR template was DNA released from bacterial colonies which had been boiled for 5 minutes. A correct clone was grown up, plasmid DNA was prepared using a Qiagen column
30 (Qiagen, Chatsworth, CA) and the DNA digested with HindIII and phosphatased.

A polynucleotide fragment containing the β Geo coding sequence was isolated from a vector provided by P.S. Mountford (Stem Cell Sciences, Melbourne, Australia). The plasmid was digested with ScaI and HindIII and the 4.3 Kb HindIII fragment containing β Geo and the polyadenylation site was gel isolated. A HindIII
5 digest alone would have resulted in two fragments of approximately equal size, therefore ScaI was added in order to digest the unwanted DNA fragment into two smaller fragments.

A ligation was performed using the pOCUS-2/PGK promoter, linearized by HindIII and phosphatased, and the HindIII β Geo fragment. Following bacterial
10 transformations, colony PCRs were carried out in order to identify a correct clone which contained the HindIII fragment in the correct orientation. The 5' PCR primer (5'AGCGCACGTCTGCCGCGCTGTT (SEQ ID NO:27)) is located in the PGK promoter, while the 3' primer (5'CCTGTAGCCAGCTTTCATCAAC (SEQ ID NO:28)) is in the β Geo fragment.

15

2.3 Construction of pPGK-EGFP

The EGFP derived plasmids were based upon the plasmid pEGFP-1 (Clontech, Palo Alto, CA; Genbank Accession #U55761) which encodes a variant of the green fluorescent protein used for monitoring the activity of promoters cloned
20 into the multiple cloning site. EGFP is human codon-optimized and contains a chromophore mutation which produces fluorescence 35 times more intense than wild-type GFP. Sequences flanking the EGFP gene have been converted to a translation initiation consensus ribosome-binding site to further increase the translation efficiency in eukaryotic cells. The vector backbone provides an SV40
25 origin of replication and polyadenylation sequence, and a neomycin-resistance cassette for selection of stably transformed mammalian cells. One construct that served as the negative control was pEGFP without further modification.

pPGK-EGP was designed to contain the mouse PGK promoter driving expression of EGFP and was used as a positive control. The vector was constructed
30 as follows: The vector pEGFP was digested with EcoRI and PstI and phosphatased. The PGK promoter was isolated by digestion of pPGKneo (Aron Thall,

BioTransplant, Inc.) using EcoRI and PstI. The two fragments, linearized pEGFP and the PGK promoter, were ligated. After transformation of *E. coli*, miniprep DNA was prepared (Qiagen, Chatsworth, CA) and correct clones were identified following digestion with EcoRI and PstI.

5

2.4 Construction of pOct-4-EGFP

The experimental construct, pOct-4-EGFP, contained the 3.2 Kb portion of the porcine Oct-4 promoter. The construct was made as follows: 5D was digested with BamHI, and the 5.5 Kb insert containing the porcine Oct-4 promoter sequence plus exon 1 and part of intron 1 was gel purified, and digested again with BstBI and FspI. The 2.84 Kb BamHI/BstBI fragment was gel purified. (The FspI was added to the digestion reaction in order to cleave an unwanted 3 kb fragment). The remaining 0.36 Kb portion of the promoter was obtained by PCR, using the 5D insert as template, and the following primers:

15

Oct4-BstBI 5' 5' CAGGGTCTTCGAAGAGGGGTCCA (SEQ ID NO:29)

Oct4-SalI 3' 5'GTCGACCAGGGCTCTCCAAGGGGA (SEQ ID NO:30)

This resulted in the addition of a SalI restriction site to the 3' end (SalI) which could be used for cloning purposes in pEGFP-1. Following PCR, the 0.36 Kb product was cloned into the TA vector (pCRII) and sequenced. After a correct clone was identified, a ligation was performed with the following DNA fragments: (1) pEGFP-1 digested with BamHI and SalI, and phosphatased; (2) the 2.84 Kb BamHI/BstBI portion of the Oct-4 promoter sequence; and (3) the 0.36 Kb BstBI/SalI 3' end of the Oct-4 promoter. Following transformation of *E. coli*, correct clones were identified by colony PCRs using the following primers:

25

5'GTCGACCAGGGCTCTCCAAGGGGA (SEQ ID NO:30)

6-T7 OL.4 5' 5'ACTTAGCACAGACACCASGACCT (SEQ ID NO:31)

A correct clone yields a 0.4 kb PCR product.

2.5 Cell culture

Mouse RW4 embryonic stem cells (ES) and mouse embryonic fibroblasts (MEFs) were obtained from Genome Systems (St. Louis, MO). The ES cells had been isolated from the inner cell mass of a 3.5 day embryo from the mouse strain 129/SvJ; the MEFs had been isolated from 14.5 day embryos. The ES cells were cultured in 80% DMEM, 15% FCS (#A-1115-L, Hyclone, Logan, UT), 2 mM L-glutamine, 0.1 mM non-essential amino acids, 10 mM HEPES, Pen/strep (100 units/ml), and 72 μ M 2-mercaptoethanol. The ES cells were maintained in the presence of murine leukemia inhibitory factor (LIF, final concentration 1000 U/ml, Gibco/BRL, Gaithersburg, MD). The MEFs were grown in 87% DMEM, 10% FCS, 2 mM L-glutamine, and 0.1 mM non-essential amino acids.

2.6 Transfections and Selection

Transfections were performed by electroporation as follows: The cells were washed twice with phosphate-buffered saline (PBS), trypsinized using Trypsin/EDTA (0.05%/0.53 mM), and centrifuged. Colonies of $0.4\text{--}1.0 \times 10^7$ cells were transfected with 25 μ g linearized DNA (sequences from sections 2.1, 2.2, 2.3 and 2.4 as described above) in 800 μ l electroporation buffer (20 mM HEPES, pH 7.05, 137 mM NaCl, 5 mM KCl, 6 mM D-glucose, 0.7 mM Na_2HPO_4) in a 0.4 cm cuvette using 300 v and 500 μ F, at room temperature. The cells were seeded in 100 mm dishes (2.5×10^6 cells/dish), containing a confluent monolayer of 30 Gy irradiated MEF. 24 hours later, the media was exchanged with selection medium containing 470 μ g/ml G418. On day 8, G418 resistant colonies were picked and transferred to a 96-well plate that contained 50 μ l trypsin solution. After 10 minutes, the cells were transferred to a 48 cell plate that contained medium plus 240 μ g/ml G418. After three days, the medium was changed to 470 μ g/ml G418 containing medium.

2.7 X-Gal (5-Bromo-4-Chloro-3-Indolyl β -D-Galactopyranoside) Staining Assay

The medium was removed from the culture dishes and the cells were washed briefly with PBS. Fixation solution (0.2% glutaraldehyde, 0.1 M phosphate buffer, pH 7.3 [4 volumes 0.1 M disodium orthophosphate + 21 volumes 0.1 M sodium dihydrogen orthophosphate], 2 mM MgCl_2 , 5 mM EGTA) was added to the cells for 5 minutes at 4°C. The cells were washed three times for 10 minutes each time with wash solution (0.1 M phosphate buffer, pH 7.3, 2 mM MgCl_2). The staining solution was added (1 mg/ml X-Gal [Sigma, St Louis MO] final concentration prepared by dissolving 25 mg X-Gal in 500 μl N,N dimethyl formamide, mixed with 25 ml of washing solution with 41 mg potassium ferricyanide and 52.5 mg potassium ferrocyanide) overnight at 37°C. The stained plates were stored in 0.2% glutaraldehyde at 4°C.

2.8 Retinoic Acid Assay

A stock solution of all trans-retinoic acid (Sigma, St. Louis, MO) was prepared by dissolving it in absolute ethanol to a final concentration of 1 mM.

The cells were seeded in 24-well plates containing a feeder layer of irradiated MEFs. After 3 days, the media was adjusted to 0.5 μM retinoic acid. The medium was changed every 2-3 days. After 5 days, the medium was removed and fresh medium without retinoic acid was added to the cells. Cells were split if necessary to avoid confluency. The colonies were then stained using the X-gal procedure.

2.9 X-Gal staining of cells transfected with the various vectors

The results (Table 1) demonstrate that the number of colonies derived from transfections using the PGK and Oct-4 promoter-containing plasmids was 50-100 fold higher than the number of colonies derived from the transfection using the promoterless plasmid. Therefore, the DNA sequence shown in Figure 8, isolated from porcine genomic DNA, contains the porcine Oct-4 promoter sequence. The single clone that resulted from the transfection using the promoterless plasmid was demonstrated to express the βGeo fusion protein. This clone proved to be useful as a

control for subsequent experiments as a non-developmentally regulated promoter. As used herein the staining legends indicate the following: XXX indicates a strong stain, XX indicates a moderate stain, and X indicates a weak stain.

5 **Table 1 Transfection Efficiency and X-Gal Staining of Neomycin Resistant**

VECTOR	TOTAL # COLONIES	X-GAL STAINING
pβGeo - no promoter control	1 (1 colony/5 x 10 ⁶ cells)	XXX
pPGKβGeo (pOCUS-2)	9 (24 colonies/10 ⁶ cells)	XXX
pOct-4-βGeo (pGT1.8)	90 (12 colonies/10 ⁶ cells)	XX

Legend: X-weak stain XX-moderate stain XXX-strong stain

2.10 X-Gal staining after treatment of cell cultures with retinoic acid

10

The results of the X-Gal staining assay using cells that had been treated with retinoic acid are shown in Table 2.

Cells grown derived from the single colony that was obtained from the pβGeo plasmid transfection differentiated in the absence of LIF. These differentiated colonies expressed the β-Geo fusion protein, further establishing that the promoter driving the expression of this protein was not regulated by the presence of retinoic acid in the culture medium.

Cells from the colonies that were derived from the pPGKβGeo transfection also underwent differentiation in the presence of retinoic acid in the culture medium, with only undifferentiated cells from one colony (A5) surviving as a mixed population of differentiated and undifferentiated cells. This result was expected due to the understanding that the PGK promoter is differentially regulated by the presence of retinoic acid (Sutherland et al. (1995) Gene Expression 4:265-279). The differentiated cells did not stain with X-Gal, indicating that indeed the PGK promoter was not active in these differentiated cells. These cells functioned in this assay as the

positive control. The cells from the pOct-4 β Geo derived colonies also differentiated in the presence of retinoic acid, with the resulting cell population being a mixture of a few undifferentiated cells together with a lot of differentiated cells. The undifferentiated cells stained with X-Gal but the differentiated cells did not. These results indicate that the porcine Oct-4 promoter is differentially regulated by the presence of retinoic acid in the culture medium. (Since retinoic acid promotes differentiation of the cells, it results in the turning off of the porcine Oct-4 promoter sequence because the sequence is only active in embryonic stem cells and not in differentiated cells.)

10

Table 2 X-Gal Staining of Colonies after Treatment of Cell Cultures with Retinoic

Vector	LIF	G418	Name of Colony	Undifferentiated colonies/well	Differentiated colonies/well	X-gal staining of undifferentiated colonies	X-gal staining of differentiated colonies
p β Geo	-	-	A2	NO	XXX	na	XX
pPGK β Geo	-	-	A2	NO	XXX	na	NO
			A3	XX	XXX	X	NO
			A5	X	XXX	X	X
pOct4 β Geo	-	-	F4/A	X	XXX	X	NO
			E3/A	NO	XXX	na	NO
			E5/A	XX	XXX	X	NO
			E8/A	X	XXX	X	NO
			B1/A	X	XXX	X	NO
			E6/A	X	XXX	X	NO
			A1/B	XX	XXX	X	NO
			D2/B	XX	XXX	X	NO
			C3/B	NO	XXX	na	NO

Legend: X-weak stain XX-moderate stain XXX-strong stain

An alternate method for inducing differentiation of ES cells is to culture them in the absence of LIF. Colonies were grown in the absence of a feeder layer but in the presence of G418 selection, for 12 days, and then subjected to the X-Gal staining protocol. The results are presented in Table 3.

The effects of the absence of LIF upon the ability of the p β Geo derived cells to undergo differentiation was less pronounced than the retinoic acid induced differentiation, but cells that did differentiate retained the ability to express the β Geo fusion protein, as expected.

Cells derived from the pPGK β Geo transfection did undergo differentiation, but did not lose the ability to express genes from the PGK promoter and therefore survived in the presence of G418. These results confirmed differences in the mechanism and/or extent of differentiation of ES cells induced by the absence of LIF but in the presence of retinoic acid or G418. The differentiated cells retained the ability to be stained by the X-Gal protocol.

Cells derived from the pOct-4 β Geo transfections were similarly able to differentiate in the absence of LIF but did not express the β Geo protein. The cells appear to be differentially sensitive to the presence of G418, since some cells were still capable of proliferation despite the down-regulation of the pig Oct-4 promoter.

Table 3 X-Gal Staining of Cells Grown in the Absence of LIF, but in the Presence of G418 Selection

Vector	LIF	G418	Name of Colony	Undifferentiated colonies/well	Differentiated colonies/well	X-gal staining of undifferentiated colonies	X-gal staining of differentiated colonies
p β Geo	-	+	A2	XXX	X	XXX	X
pPGK β Geo	-	+	A2	X	XXX	X	X
	-	+	A3	X	XX	XX	X
			A5	XX	XX	X	X
pOct4 β Geo	-	+	F4/A	X	XX	X	NO
	-	+	E3/A	X	XX	X	NO
	-	+	E5/A	X	XX	X	NO
	-	+	E8/A	X	XX	X	NO
	-	+	B1/A	XXX	XX	XXX	NO
	-	+	E6/A	XX	XX	XX	NO
	-	+	A1/B	XX	XX	X	NO
	-	+	D2/B	XX	XX	XX	NO
	-	+	C3/B	X	X	X	NO

Legend: X-weak stain XX-moderate stain XXX-strong stain

2.11 Transfection of porcine Oct-4/EGFP in mouse ES cells

Linearized plasmid DNAs were co-transfected with pPGKneo in a 1(PGK-Neo):10 (pOct-4/EGFP) ratio. Cells were selected for the ability to grow in 470 μ g/ml G418. On day 7 fluorescent positive colonies were picked. No colonies grew from the promoterless control transfection; six colonies grew from the pPGK-EGFP

transfection, representing six colonies/million cells plated; eleven colonies grew from the pOct-4-EGFP transfection, representing 3 colonies/million cells plated. Colonies were subjected to the retinoic acid differentiation protocol, namely the colonies were grown in 0.5 μ M retinoic acid for 5 days, then 7 days without retinoic acid (growing on MEFs). After this treatment all of the colonies lost the ability to fluoresce, thus confirming that the pig Oct-4 promoter is developmentally regulated and hence is an ES cell-specific promoter.

As described above, Applicants' have provided the porcine Oct-4 promoter polynucleotide sequence which enables one to selectively propagate large numbers of porcine embryonic stem cells. Once this is accomplished, one skilled in the art is then able to create a transgenic pig line which will comprise the cells which will express the selectable marker under the control of the porcine Oct-4 promoter sequence. Technology to create such a transgenic animal is known in the art and can be found, for example, in Transgenic Animal Technology, A Laboratory Handbook ((1994) ed., Carl A. Pinkert, Academic Press, Inc., San Diego, CA).

Additionally, one skilled in the art may propagate large numbers of stem cells by utilizing the selectable marker under the control of the porcine Oct-4 promoter polynucleotide sequence as described above, and then further genetically modify these cells to accomplish a desired activity or to eliminate an activity (e.g., to eliminate α 1,3 galactosyltransferase activity). (See, e.g., Tearle et al. (1996) Transplantation 61:13-19 entitled "The α 1,3 galactosyltransferase Knockout Mouse: Implications For Xenotransplantation."). These cells that have been further genetically modified are then capable of being used to generate a transgenic pig line containing cells, and organs, with the desired genetic activity or eliminated activity.

Although the invention has been described with reference to its preferred embodiments, other embodiments, can achieve the same results. Variations and modifications to the present invention will be obvious to those skilled in the art and it is intended to cover in the appended claims all such modification and equivalents and follow in the true spirit and scope of this invention.

We claim:

1. A method of isolating or selectively propagating porcine stem cells, wherein said method comprises introducing into a source of cells containing porcine stem cells a genetic selectable marker construct which is operatively linked to a porcine promoter polynucleotide sequence which provides differential expression of the selectable marker in stem cells and cells other than the desired stem cells, and which under appropriate culture conditions enables the selective isolation and/or propagation of the desired stem cells.
2. The method of claim 1 wherein said porcine stem cells are embryonic stem cells.
3. The method of claim 1 wherein said porcine stem cells are pluripotential stem cells.
4. The method of claim 1 wherein the promoter polynucleotide sequence comprises a porcine Oct-4 promoter polynucleotide sequence.
5. The method of claim 4 wherein the porcine Oct-4 promoter polynucleotide sequence comprises the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
6. The method of claim 4 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence having at least 90% sequence identity with all or a contiguous portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
7. The method of claim 4 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence having at least 95% sequence identity with all or a contiguous portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).

8. The method of claim 4 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence having at least 98% sequence identity with all or a contiguous portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
9. The method of claim 4 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence which hybridizes under high stringency conditions to all or a contiguous portion of the Oct-4 promoter polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
10. A method of selectively isolating and/or propagating porcine stem cells, said method comprising culturing a source of cells under selective culture conditions, characterized in that the source of cells includes porcine stem cells containing a genetic selectable marker whereby a gene product associated with the genetic selectable marker is produced and which under said culture conditions causes selective reproduction of the desired stem cells to occur, and wherein the genetic selectable marker is operatively linked to a porcine promoter polynucleotide sequence that regulates expression, which promoter polynucleotide sequence is differentially active in stem and non-stem cells.
11. The method of claim 10 wherein the promoter polynucleotide sequence comprises a porcine Oct-4 promoter polynucleotide sequence.
12. The method of claim 11 wherein the porcine Oct-4 promoter polynucleotide sequence comprises the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
13. The method of claim 11 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence having at least 90% sequence identity with all or a contiguous portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).

14. The method of claim 11 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence having at least 95% sequence identity with all or a contiguous portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
15. The method of claim 11 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence having at least 98% sequence identity with all or a contiguous portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
16. The method of claim 11 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence which hybridizes under high stringency conditions to all or a contiguous portion of the Oct-4 promoter polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
17. An Oct-4 promoter polynucleotide sequence comprising the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
18. A polynucleotide sequence capable of hybridizing under conditions of high stringency to all or a portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
19. A porcine stem cell(s) capable of being cultured under appropriate selective culture conditions so as to enable isolation and/or propagation of porcine stem cells, characterized in that said porcine stem cell(s) contains a genetic selectable marker wherein differential expression of the selectable marker in the desired stem cell(s) and cells other than the desired stem cell(s) enables selective survival or growth of the desired stem cell(s) to occur, and wherein said genetic selectable marker is operatively linked to a porcine promoter polynucleotide sequence which provides differential expression of the selectable marker in stem cell(s) and cells other than the desired stem cell(s).

20. The porcine cell(s) of claim 19 wherein the porcine promoter polynucleotide sequence comprises a porcine Oct-4 promoter polynucleotide sequence.
21. The porcine cell(s) of claim 20 wherein the porcine Oct-4 promoter polynucleotide sequence comprises the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
22. The porcine cell(s) of claim 20 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence having at least 90% sequence identity with all or a contiguous portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
23. The porcine cell(s) of claim 20 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence having at least 95% sequence identity with all or a contiguous portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
24. The porcine cell(s) of claim 20 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence having at least 98% sequence identity with all or a contiguous portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
25. The porcine cell(s) of claim 20 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence which hybridizes under high stringency conditions to all or a contiguous portion of the Oct-4 promoter polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
26. A transgenic pig which comprises a source of porcine cells suitable for the isolation and/or propagation of stem cells by a method according to any of claims 10 to 16.

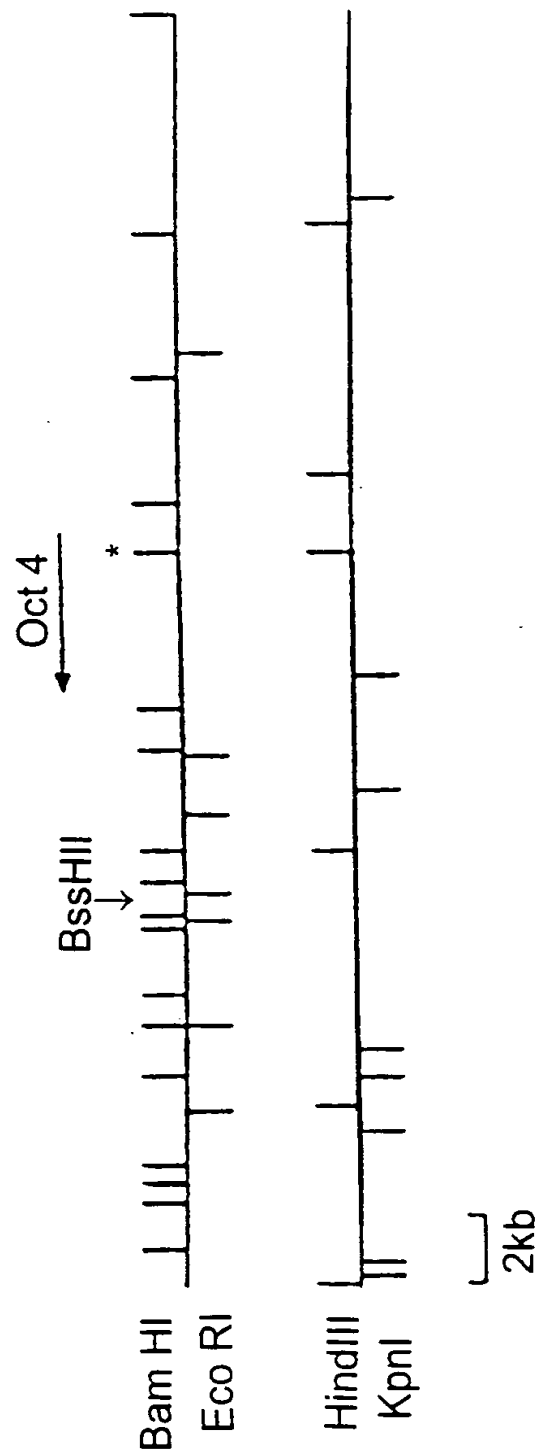
27. A transgenic pig generated using a cell obtained by a method claimed in any of claims 1 to 9.
28. The transgenic pig according to claim 27 comprising cells which include said genetic selectable marker.
29. A vector for use in genetically modifying cells so as to be suitable for use as the source of cells defined in any of claims 1 to 16, said vector comprising a first genetic component corresponding to said genetic selectable marker and a second genetic component which in the genetically modified animal cells directly or indirectly results in the said differential expression of the selectable marker, and wherein said second genetic component comprises a porcine promoter polynucleotide sequence which is differentially activated in stem cells and cells other than stem cells.
30. The vector of claim 29 wherein said porcine promoter polynucleotide sequence comprises a porcine Oct-4 promoter polynucleotide sequence.
31. The vector of claim 30 wherein said porcine Oct-4 promoter polynucleotide sequence comprises the nucleotide polysequence as shown in Figure 8 (SEQ ID NO:24).
32. The vector of claim 30 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence having at least 90% sequence identity with all or a contiguous portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
33. The vector of claim 32 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence having at least 95% sequence identity with all or a contiguous portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).

34. The vector of claim 32 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence having at least 98% sequence identity with all or a contiguous portion of the polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
35. The vector of claim 32 wherein the porcine Oct-4 promoter polynucleotide sequence comprises a polynucleotide sequence which hybridizes under high stringency conditions to all or a contiguous portion of the Oct-4 promoter polynucleotide sequence as shown in Figure 8 (SEQ ID NO:24).
36. The vector of claim 29 wherein the genetic selectable marker is an antibiotic marker.
37. The vector of claim 31 wherein the genetic selectable marker is an antibiotic marker.
38. A method of preparing a transgenic pig, said method comprising the steps of
 - providing a blastocyst;
 - providing porcine cells according to any of claims 19 to 25;
 - introducing the porcine cells into said blastocyst;
 - transferring the blastocyst to a recipient; and
 - allowing an embryo to develop to an animal to enable germline transmission of the selectable marker.
39. The porcine cell(s) of any of claims 19 to 25, said porcine cell(s) further comprising a genetic manipulation to eliminate $\alpha 1,3$ galactosyltransferase activity of said cell(s).

40. A transgenic pig comprising a porcine cell(s) according to any of claims 19 to 25.
41. A transgenic pig comprising a porcine cell(s) according to claim 39.
42. An organ suitable for transplantation into a recipient, said organ taken from a transgenic pig according to claim 41.

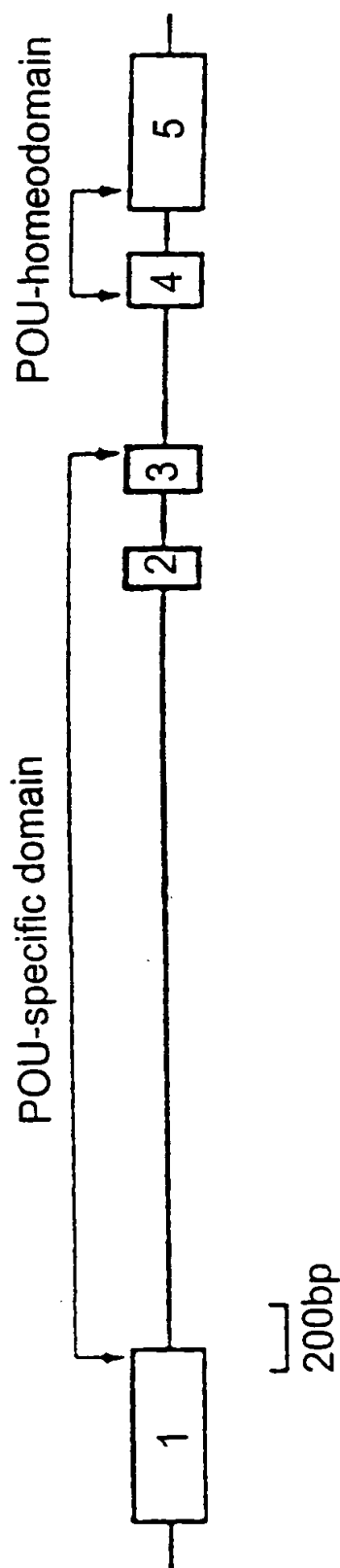
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FIG. 1A




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FIG. 1B



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FIG. 2A

BamHI SmaI PstI EcoRI 

AGTGGATCCCCCGGGCTGCAGGAATTCGGG

BamHI

ATGGATCCTCGAACCTGGCTAAGCTTCCAAGGGCCTCCAGGTGGGCTGGAATCGGACCA 60
 1 M D P R T W L S F Q G P P G G P, G I G P -


GGCTCAGAGGTATTGGGGATCTCCCCATGTCCGCCCCGATACGAGTTCTGCGGAGGGATG 120
 21 G S E V L G I S P C P P A Y E F C G G H -


GCATACTGTGGACCTCAGGTTGGTCTGGGCTAGTCCCCCAAGTTGGCGTGGAGACTTG 180
 41 A Y C G P Q V G L G L V P Q V G V E T L -

CAGCCTGAGGGCCAGGACGAGTGGAAAGCAACTCAGAGGGNACCTCCTCTGAG 240
 61 Q P E G Q A G A R V E S N S E G T S S E -

CCCTGTGCGGACCGCCCAATGCCGTGAAGTTGGAGAAGGTGGAACCAACTCCCGAGGAG 300
 81 P C A D R P N A V K L E K V E P T P E E -

TCCCAGGACATGAAAGCCCTGCAGAGGAGCTAGAACAGTTTGCCCAAGCTGTGAAGCAG 360
 101 S Q D H K A L Q K E L E Q F A K L L K Q -

PstI  Clone 1

PstI  Clone 16

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FIG. 2B

_____POU-SPECIFIC DOMAIN _____
 AAGAGGATCACCTTGGGGTACACCCAGCGGACGTGGGGCTCACCCCTGGGCGTCTCTTT 420
 121 K R I T L G Y T Q A D V G L T L G V L F -

 GGAAGGTGTTCAAGCCAGACCACTCTGTGCTTCGAGGCGCTTGCAGCTCAGCCTTAAG 480
 141 G K V F S Q T T I C R F E A L Q L S L K -

 AACATGTGTAAGCTGCGGCCCTGCTGGAGAAGTGGGTGGAGGAGCCGACAAACAATGAG 540
 161 N H C K L R P L L E K H V E E A D N N E -

 AACCTTCAGGAGATATGCAAAATCGGAGACCCCTGGTGCAGGCCCGGAGAGAAAGCGAACT 600
 181 N L Q E I C K S E T L V Q A R K R K R T -
 _____POU HOMEODOMAIN _____
 AGCATTGAGAACCGTGTGAGGTGGAGTCTGGAGACCATGTTTCTGAAGTGCCCGAAGCCC 660
 201 S I E N R V R W S L E T H F L K C P K P -

 TCCCTACAGCAGATCACTCACATCGCCCAATCAGCTTGGGCTAGAGAGGATGTGGTTCGA 720
 221 S L Q Q I T H I A N Q L G L E K D V V R -

 GTATGGTTCTGTAAACCGCGCCAGAGGGCAAAAGATCAAGTATTGAGTATTCCTCAACGA 780
 241 V H F C N R R Q K G K R S S I E Y S Q R -

FIG. 2C

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GAAGAGTATGAGGCTACAGGGACACACCTTTCCACGGGGGGCTGTATCCTTTCCTCTGCCC 840
 261 E E Y E A T G T P F P G G A V S F P L P -

 CCAGGTCCCCACTTTGGCACCCAGGCTATGGAAGCCCCCACTTCACCACTCTACTCA 900
 281 P G P H F G T P G Y G S P H F T T L Y S -

 GTCCCTTTTCCCTGAGGGCGAGGCCTTTCCCTCTGTTCCTGTCACCTGCTGGGCTCTCCC 960
 301 V P F P E G E A F P S V P V T A L G S P -

 ATGCATTCAAACTGAGGCACCAAGCCCTCCCTGGGGATGCTGTGAGCCAAGGAGGAGG 1020
 321 H H S N * 324

 TAGACAAGAGAACCCTGGAGCTTTGGGGTTAAATTCTTTTACTGAGGAGGGATTAAAGCA 1080
 CAACAGGGGTGGGGGTGGGATGGGAAGAAGCTCAGTGATGCTGTCAGGAGCCT 1140
 GGCCTGCTGTCACTCATTTTGTCTTAATAAAGACTGGGACACACAGTAAAAAA 1200
 AAAAAAAAAAACTCGAG 1217
 XhoI

FIG. 3A

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10	20	30	40	50
* * *	* *	* *	* *	* *
GAATTCGGCT	TCACCAGGCT	TCGGACTTCG	CCTTCTCGCC	CCCGCCGGGC
60	70	80	90	100
* * *	* *	* *	* *	* *
GGTGGAGGCG	ATGGGCCCGG	AGGGCGGAGC	CGGGCTGGGT	TGATCCTCGG
110	120	130	140	150
* * *	* *	* *	* *	* *
ACCTGGCTGA	GCTTCCAAGG	GCCTCCCGGT	GGTCAAGGA	TCGGGCCGGG
160	170	180	190	200
* * *	* *	* *	* *	* *
GGTTGGGCCG	GGCGCCGAGG	TGTGGGGGCT	TCCCGCGTGC	CCCCCGCCCT

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FIG. 3B

210	220	230	240	250
* * *	* * *	* * *	* * *	* * *
ATGACTTCTG	CGGAGGGATG	GCCTACTGCG	CACCTCAGGT	CGGAGTGGGG
260	270	280	290	300
* * *	* * *	* * *	* * *	* * *
CTGGTGCCCC	AGGGCGGCCT	GGAGACCCCT	CAGCCCCGAGG	GCGAGGCCGG
310	320	330	340	
* * *	* * *	* * *	* * *	
GGCCGGGGTG	GAGAGCAACT	CCGAGGGGGC	CTCCCCCGAG	

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FIG. 4A

10	20	30	40	50
* *	* *	* *	* *	* *
GGATCCTCGG	ACCTGGCTGA	GCTTCCAAGG	GCCTCCCCGGT	GGGTCAGGGA
60	70	80	90	100
* *	* *	* *	* *	* *
TCGGGCCCGG	GGTTGGGCCG	GGCGCCGAGG	TGTGGGGGCT	TCCCCCGGTGC
110	120	130	140	150
* *	* *	* *	* *	* *
CCCCCGCCCT	ATGACTTCTG	CGGAGGGATG	GCCTACTGCG	CACCTCAGGT
160	170	180	190	200
* *	* *	* *	* *	* *
CGGAGTGGGG	CTGGTGCCCC	AGGGCGGCCT	GGAGACCCCT	CAGCCCCGAGG

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FIG. 4B

210 220 230 240 250
* * * * *
GCGAGGCGGG GCGCCGGGTG GAGAGCAACT CCGAGGGGGC CTCCCCCGAG

260 270 280 290 300
* * * * *
CCCTGTGCCG CCCCCGCTGG CGCCGCGAAG CTGGACAAGG AGAAGCTGGA

310 320 330 340 350
* * * * *
GCCGAACCCC GAAGAGGCCA GTGAGCTGCC GGGAGCTGGG GGAGGCGATC

360 370 380 390 400
* * * * *
GCGCTGGCCG GGGCGGCACG CAGGGAGGT GGTCCCTGC CGCCCGGGCA

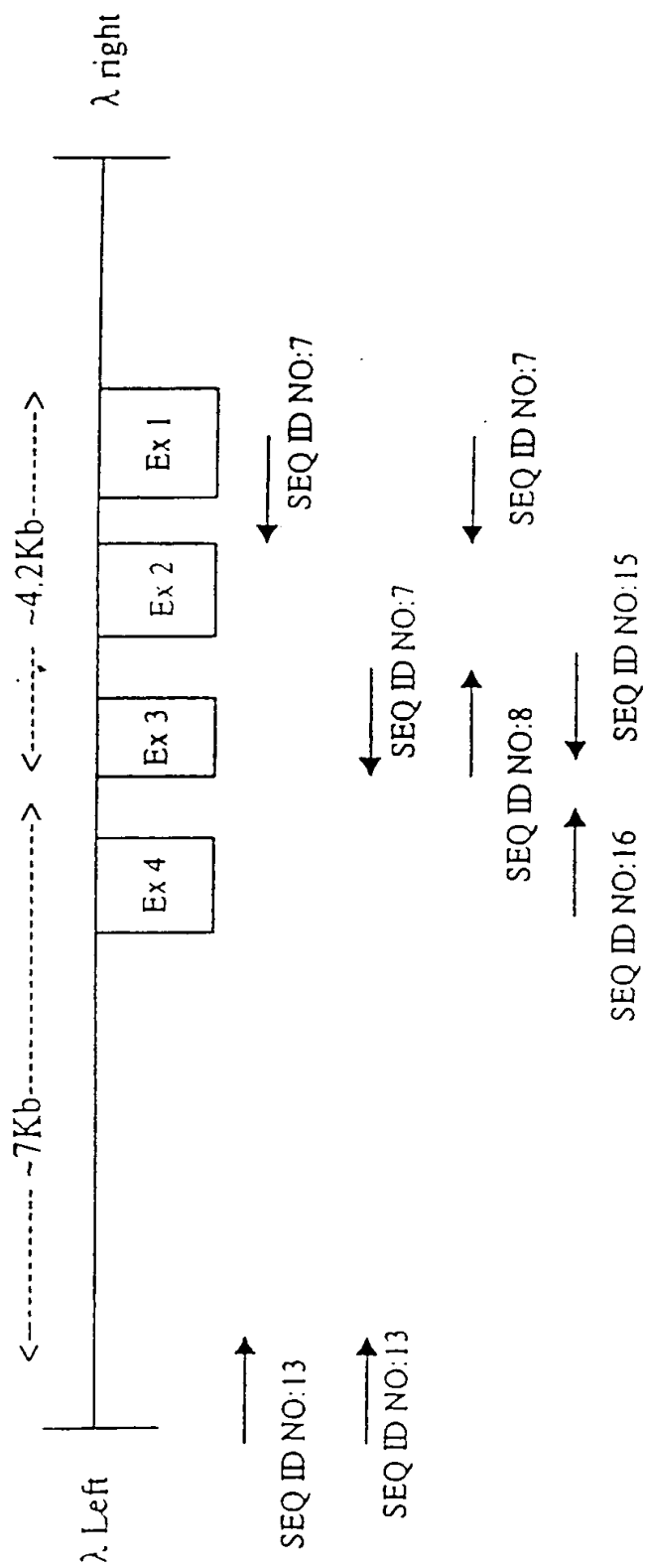
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GGAGGGG

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FIG. 5A

Oct-4 CLONE 3



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FIG. 5B

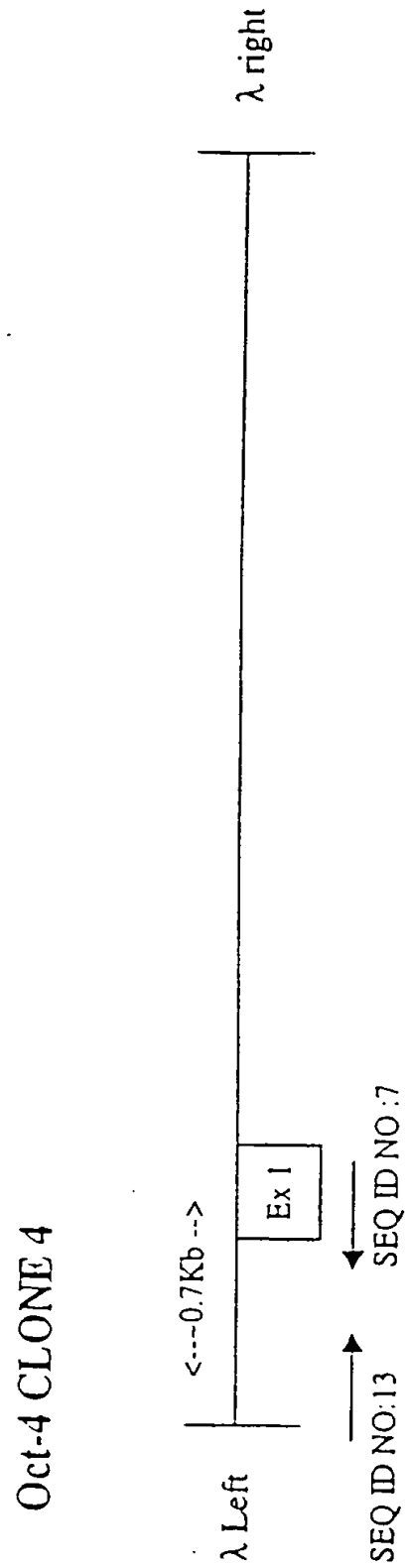


FIG. 6A

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10	20	30	40	50
* * *	* *	* *	* *	* *
AGCCATGCTG	GGTGATCCT	CCCACCTGGC	TGTGCTTCCA	AGGGCCTCCT
60	70	80	90	100
* * *	* *	* *	* *	* *
GTTGGGTCAG	GGATCTGGCG	GGGGGCTGCT	GGACCCAGAG	GTGGGGAGGC
110	120	130	140	150
* * *	* *	* *	* *	* *
TTCTCTCATG	CCCCCCGCCC	TAGGACTTCT	GCGGAGGGAT	GGCCTACTGT
160	170	180	190	200
* * *	* *	* *	* *	* *
GCACCTCAGG	TCAGAGAGGG	GCTGGTGCCC	CAAGGCGGCC	TGGAGACCCC

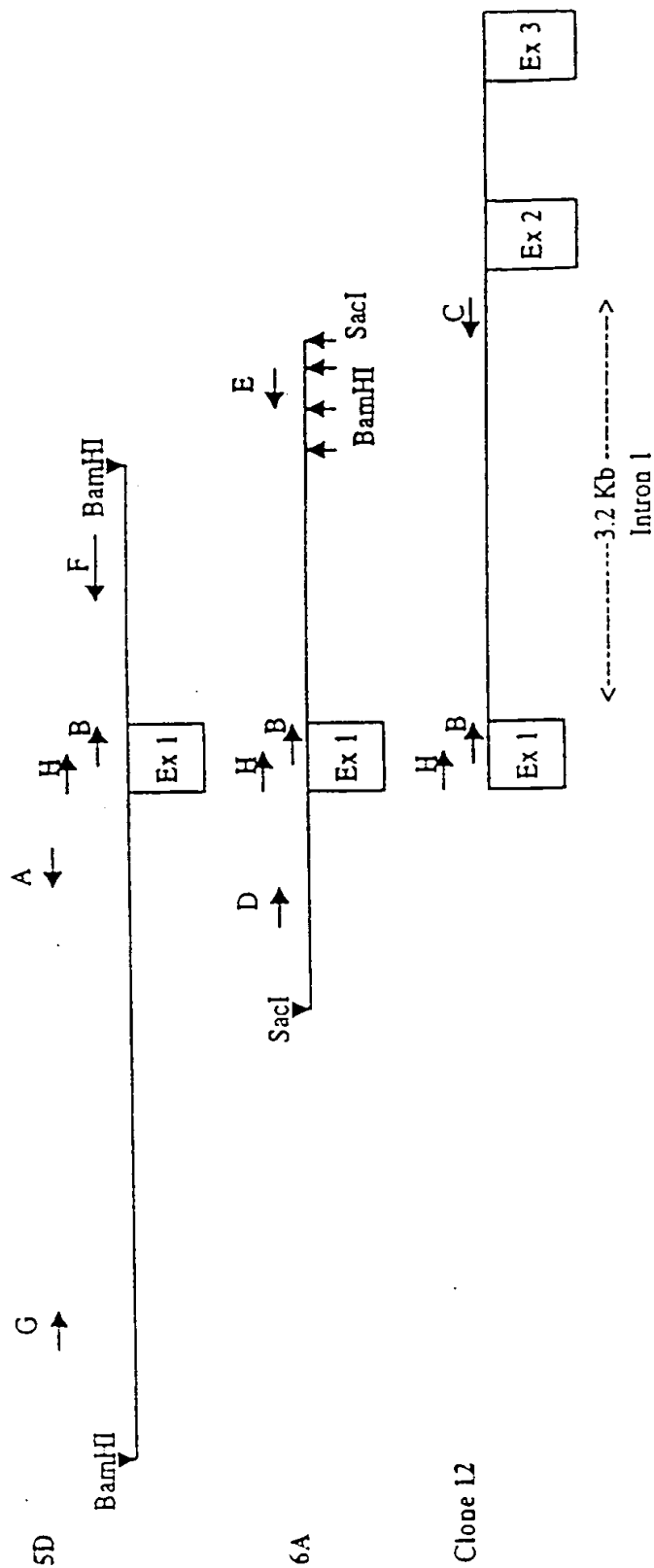
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FIG. 6B

210	220	230	240	250
* *	* *	* *	* *	* *
TCAGCCCTAG	GGCCAGGCAG	GAGTCGGGGT	GGGAGCAAC	TCCGAGGGGG
260	270	280	290	300
* *	* *	* *	* *	* *
CCTCCCCTGGA	GCCCTATGCC	ACCCCCGTTG	GCACTGCACA	GCTGGACAAG
310	320	330	340	350
* *	* *	* *	* *	* *
GAGAAACTAG	AGCCGAATCC	TGAGAAGTCC	CAGGACATCA	AAACGCTTCA
360	370			
* *	* *			
GAAAGACCTT	CAACAATTG	CCAAGCTT		

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FIG. 7



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FIG. 8A

GGATCCCTAG	CCTGGAAACC	TCCATAAGCC	GTGGGTACTG	CCCTAAGAAG	AAAAAAAAG	60
TGGTTTGCTA	CCCTGGTCTA	GAGCAAGCCT	CCATTTTCCC	CAGGAGTCAT	TTCAGCTGGT	120
TTTCCCTACC	AAACAGCAAG	GGGATGGCCG	GGCTGACAGC	AGCAAAGTCA	CTGTCACTTC	180
TTTGCAGCCT	TGCCAGGCCA	GCTGCATCTG	GCAGGAGCG	GCAGCTCTCA	CCTGCCCTCC	240
CTGGGTCATG	CTCTACAGCC	AGATACTTGG	CATTTGTCTT	TGTGTAGGGC	CTCAATATTG	300
TACTCTAATA	AGGTACATG	TGGGAGTTCC	CTGGTGGCTC	AGCAGTTGAG	GATCTGGCAT	360
TGTCACTGCT	CTGGCATGGA	TCTCTGCTCT	GGCGCAGGTT	CAATTTCCTGG	CCTGGGAACT	420
TCTGTACGCC	GCAGGCGTGA	CTGGAAAAAT	ACAGGTGGGG	TGGGGTGAGG	AGTGTATGTG	480
GAGAGTCTGC	AAACCCAGGC	CTAAATTGGT	TTGGGGGACT	TGAAGTTTTT	AGTGACTCCC	540
TACCCAAAAG	AGTGAGAGAAG	CCAGGTCTGA	TGACTTAAACC	CCACTTGCCAG	TCTGCTCTGG	600
GCCTGCAGAG	ACCTGGCCTC	TGCAGAAAGT	AAGCTGCCCTA	CAC TTCAGGC	CTAACAGGAG	660
GGTGGGGAGG	AGAGGGGAAT	AGGCTCAGCC	CTGCCATGCC	AAGCACCCCC	AGGCTGACTA	720
GGACTCCAGA	CAAA'TTTAGC	TTGTCCCTTAA	GGTTCCTGGGT	CAGACCCCCAG	GCAAGCACAG	780
AACTGATCTG	GCTCAGATGT	CTGGCTACAA	GCTATCCAGG	AACCCAGGCA	TCCAGCCCTC	840
CCCAGCCCTC	CCCAGGCTTT	CCCTCTGGAA	TAGGAAGGAC	ACTTGCTTAA	ACCAGAAACA	900
TACCATCTAG	AGCAGCTATT	TATGGTGATC	TAAAAAACAC	AGGGTGCTAT	TTAGTCGGGG	960
GTGGGTGGGA	AGGGAGAAGG	TGTTTAGGGT	CCGCGGGAAG	GTCAGGGGCA	CAGGGGCTCT	1020
CTGGACCACA	TGGGGAGAGG	GGTTTCTGGG	AGGCCAGAGG	GCGAGGAGCC	AAGGAGCTCA	1080

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FIG. 8B

GCAGTAGATT CCCTAGGCC GCCCCTCCCC CTCCTCAGGG AGGCCGTCTT CTGGCAGAC 1140
 AGCAGATAGA TGCATGACAA AGGGCCCATG ATGGCTCTGT CCTGGGGGTT GGGGAGATGG 1200
 CTAGGGAGGG GCCCCTCCTG GTCTGAAGCA CATCTTTCCA CCCCACCAG GCCCCTTAAT 1260
 CTATCTGCTT TTGGGGCAGT TAGTAGTTTA GAGTTGAAAT AGCTCCAGCC CTGCTGCCCT 1320
 ATAAATCTTT CAACAGACCT ATGGGAAGTA TTGAAATGCA TGCACGCAAT TAGTCACCCC 1380
 AAACGCACAG GCCGATGGG ACTGGAAGAG ATTCAGAGGA GAAAAAGCAA AACAAAACAA 1440
 CACGGACACA CAAAAACCCA ACAGACTCAA AGGACTCCTG GTGGAGCTAA CTGGTCACAG 1500
 TCTGGAGGAT GCCAGCCCCCT CAAGACAGAT GCCGAGCCAC TGACCCCTAGC AAACAACCTC 1560
 AGACCCAGCC AAGATGAAGA GGTGTCTAGG TCCGCAGAGG TCTGTGTCCC AGTCTCAGGA 1620
 GTCTGGCCTC CAAACTGTAG GAAGCTCTGA TCCATGGCTT CTCTGGAGAG CCCCCTCAC 1680
 TCAGGTTTAC CTGGGGCCTT CGTTTAGGGC AAGTTGGGG AGCAGACAGA CAAACATCAT 1740
 CCCCAGCAGA CAGCCAGTCT GAAAGCTATT CTCTTGCAA CAGAATCAAG CACTAGGCCA 1800
 GCAGCCTGAG CCTCAGGACA GACCCAGAAA AATAGACCCT GTGGGAGAGC TTAGGGCAGG 1860
 ATTCCCTGCAC CCCCTCCCCA ATCGCAGTTC ACCCCCTTCT GCATCTTTTC GCTAGCCCCC 1920
 CAAACAAAGG CCTGGACGCC TCAGTCCTCT AGAGGGGGA CAGGATACCT AGTCCCAGT 1980
 GGGGGGCCCT GTCTGAGGCT CAGTCTTTGA GGGGATGGGG GTGTTGTTGC TGGAGCTCTT 2040
 TTAGCTGCTC TGAAGGGAT TCTGTGTGAG GGGATTGGGG CTGGGGGGTT GGGGGGCAGG 2100
 AAGCTGTCCC CAGGGGAGCC ATCCAGGCC ATTCAAGGT TGAGCCTTG TTAGGGTTA 2160

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FIG. 8C

GAGCTGCCCC CTCGTGGGAC CAGGATTGTC CAGCCAAGGC CATTGTCCGG CCCCCTTCCC 2220
 CCAGTCCCTC CCAAGCCTCT TTGAACCTGA AGTCAGATAT TTTTCTCTCC CCCCCCTCC 2280
 CTCCTTGGCT TTCCCCACCC AGGCCCTAGG GGTGGAGGCC CAGATTGGGA GGTGGGGAG 2340
 GGAGAACAGT CAACTATGG GCTAGATAAT TGGTCCCTG AAGGGGGCT GGGGGACAAG 2400
 GAACCTGATG TCGCGGGGA CCCACAGCG GGGACCTGCG AGCGGGTGC CGATTGATC 2460
 CTCTGCCCTGC ACAAAGATTG GGAGACTCAG GCCCAGTCCA TCGAGCTTGA TCCCTGGAAG 2520
 GGAATAATGG GGTTCATCC CTGGGTCTGG TGAAGGGAG GCCCCGGAAC CCGGAAAAC 2580
 GTACGGAATG GAAGCCCGTG TGGCAGTCTG CCCCTGGTGA GGGGTGGAAT CTAATAGGCT 2640
 GGGCGGATGG TTGCTGGCA TCGCAGCTTT GGGGTGCCCG AATCTGGCCA GTAATCTAGT 2700
 TGGGAATGCC TAGGTTCCCG GACTGGGGT GAGGCGAGAG AGCAGGAAT GAGGAGTAGC 2760
 TCCGGCAGGA CTTAGCACAG ACACCAGACC TGTGTGAGGA CCTGAGAGGG TCGCTGGGGT 2820
 CCTTTGAGGA GACAGTGCCA GGGTCTTCGA AGAGGGGTCC AACACCTGGC TCCCCGACAG 2880
 CCCCAATGTG CACAGAGCAG TGGAGAGGC CGGCGGCCG GTTGGAGTT GGAGGTGAAG 2940
 GCCGCATGGG GGACCTGCAC CAAGGGCCTG GGGACCCGAG AGGCGCCCG GCGACCTCT 3000
 CCGACTTTCG CCTCCAGAC ACCACCGCCA CCAGCCAGCA AACACCTCC GCCTCAGTTT 3060
 CTCCCACCCC CACCGACCCC TCCCCACCC ATCCAGGGG CGGGGCCAGA GGTCAAGGCT 3120
 AGTGGGTGGG ATTGGGGAGG GAGAGAGGTG TCGAGCAGTC CCCTTGGAGA GCCCTGGTTT 3180
 TACTGGGCC CCGGCTGGG GCGCCTTCCT TCCCC 3215

pig oct4 2701-3215	6
huOCT4 1-499	
mOCT4 1401-1950	50
<p>-----TGGGAA</p> <p>TGCAATGGCT GTCTTGTCCT GGCCTTGGAC ATGGGCTGAA ATACTGGGTT</p>	
pig oct4 2701-3215	53
huOCT4 1-499	42
mOCT4 1401-1950	95
<p>TGCCTAGGTT CCCGACATGG GGGTGGGEC AGACAGCAG GAATTTTAG</p> <p>-----GAPT TCATGACC-A GCCTGGGTA ACATAGCTAG GCCCPTCTC</p> <p>CACCCATA-T CTAGGACCTCT AGACGGGIG GGTAAGCTA-GAACTTAG</p>	
pig oct4 2701-3215	95
huOCT4 1-499	91
mOCT4 1401-1950	140
<p>CACTAGCTCC GGCAGACTT -TA--GCACA G-ACACCAG ACCCTGTGT</p> <p>TATTAATAAT AATAAACAATA ACAGGGCACA GTGGTCCAA GCCTGTATCTC</p> <p>CACTGGCCCC AGAAA--TA AT-TGGCACA GACACATTCA A-TGATGTIT</p>	
pig oct4 2701-3215	138
huOCT4 1-499	139
mOCT4 1401-1950	183
<p>-CAG-GACTT GAGAGG--GT CGCTGG--GGT TCCCTTTTAGG AGTCAGTTC</p> <p>CCAGCCACTT AGGAGGCTGG AGCTGACGA TTGCTTTTGG -CCAGTATGA</p> <p>TTAGCTCTCTC CAGAGGATG- -GCTGATGG GC-TCTTAGG A--CAG--CTC</p>	

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FIG. 9B

pig oct4 2701-3215	C-AGGTCCTT	GGAAGAGGG	TCCACA--C	CTGGCTCC	GACAGCC	185
huOCT4 1-499	T-CGCGCTTA	CATTAG--	-CCAACA--T	TGTACTCCAC	TGCA-CTCCA	181
moOCT4 1401-1950	CGCGCGCTTA	C---AGIG-	-CCAFACGGC	TTTGTGCTCC	ATATGGGG--	225
pig oct4 2701-3215	ATGTGACAG	AGCAGTGGG	AGG-GCCGG	CGGCCGGTTG	GGAATTGGAG	234
huOCT4 1-499	GTCTGGGCA	CAAGTTCG	A---CCCTCT	C--TTAAAAA	ATATAAATAA	225
moOCT4 1401-1950	ATCCGAGCA	CTGGTTCG	AGGTGTCCG	TGACCCA-AG	GCAATGGGTGA	274
pig oct4 2701-3215	GTGAAGGCG	CATCGGAC	CTGACCAAG	GGCTGGGGA	CCGCAGAGG	284
huOCT4 1-499	AAATAGTTTC	TGTGGGAC	CTGACTGAG	GTCTCTGG--	-----AGGGG	268
moOCT4 1401-1950	GAGCACCTTG	AAGGTTCGA	ATG-ARGGCC	TTCTCTGGGT	CC-----C	316
pig oct4 2701-3215	CTCTGGCGG	ACCTCTCGA	CTTTCGCTT	C--CAACAC	CACCGTACC	332
huOCT4 1-499	CTC-AGTCTG	GTCTC-CCG	TTTTCCTCT	CTACACAC	CATTCTACC	316
moOCT4 1401-1950	GTCC-TAATG	GT-TTCTCTG	-TTCCAGCTT	CTCCA---AC	CTCCGTTCTGG	360
pig oct4 2701-3215	ATC---GAG	CATACAC-CC	TCTCGCTCAG	TTTCTCTCCAC	CCCACCTGAC	377
huOCT4 1-499	ACCAT-TAAG	CATACATCC	TTTCCTCAG	TTTCTCTCC--	-CCCTCTCTCC	362
moOCT4 1401-1950	ATGACACAG	CATATTA-GG	CTCGCTCAG	TTTCTCTCCACC	CCCACAGCT	408

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FIG. 9C

A			
pig. oct4 2701-3215	CC-CTCC	CC ACCCA	CCAG GGGCGGGC CAGAGGTCAA GGCTAG
huOCT4 1-499	CT-CTCC	CC ACCCA	CCAG GGGCGGGC CAGAGGTCAA GGCTAG
mOCT4 1401-1950	CT-CTCC	CC ACCCA	CCAG GGGCGGGC CAGAGGTCAA GGCTAG
			426 411 458
pig oct4 2701-3215	TGGGA	TGGG GAGGGAGAGA	GGTGTCCAGC GTCCCTTTG BACAGCC
huOCT4 1-499	TGGGA	TGGG GAGGGAGAGA	GGGTTCAGT GTCCCTTTG CA-AGCC
mOCT4 1401-1950	TGGGA	TGGG GAGGGAGAG-	---GTGATAC GTCCCTTTG GTGAGCC
			476 460 503
pig oct4 2701-3215	GTATAC	TGG GCGCCCGGCT	TGGGGGCGCT TCCTTCCCC
huOCT4 1-499	ATATAC	TGG GCGCCCGGCT	TGGGGGCGCT TCCTTCCCC
mOCT4 1401-1950	TTTCCAC	TGG GCGCCCGGCT	TGGGGGCGCT TCCTTCCCCA TGGCTGG
			515 499 550

TCACAGTCTG	GAGGATGCCA	GCCCCCTCAAG	ACAGATGCCG	AGCCACTGAC	1544
-----	-----	-----	-----	-----G	440

CCTAGG	AAC	AAC	CTCAGAC	CCAGCC	CAACA	TGACGACG	TC	AGGTGG	1594
CCTAGG	AAC	A-	GCTCAG-	-	-	GC-	-	GCTGG	475

CAGAGCTCTG TGTCCGCGC TTAGGAGTCT GGCCTCCATA CTGTGGAGG 1644
 CAGAGACTCTG T---CGGCG T-----A-----TATA CTGTGGGTG 510

CCTCTGATCCA	TGGCTTTCTT	GGAGAGCC	CCCTTAATCAG	GTTC-CCCTGG	1694
CCTCTG-----	-GGCTTTTCT	---GAGCCCG	TGCTTAATCA-	---C-CCCTGG	545

GGCCTTCGTT	TACGGCAGGT	TGGGGAGCA	GACAGACAAA	CATCATCCCT	1744
GGCCTTCGTT	TACGGCAGGT	TGGGGAGCA	GACAGACAAA	CATCATCCCT	595

AGCAGACAGC	CACTCTGABA	GCTATTCTCT	TGCAAAACAA	ATA	ATCAAGCACCT	1794
TGCAGACAGG	CACTCTGABG	GCTATTCTCT	TGCAAAACAA	ATA	ACTAAGCACCC	645

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FIG. 10B

Pig Oct-4 promoter Mus S58422S1	AGGCCAGTGG CCTGAGCCCTC AGGACAGGCC CAGAAAATA GACCCCTCTTGG AGGCCAGTAA TGGGATCCCTC AGACTGGGCC CAGAAAACCA ----CTTTTGG	1844 691
Pig Oct-4 promoter Mus S58422S1	GACAGTTTAG GGAGGTTTC CTGCACCCCC TOCCCAATCG CAATTTCACCC GATAGTTTAG GGAGGTTCT CTGCACCCCC TOC ----TCC TATTTCCGTC	1894 737
Pig Oct-4 promoter Mus S58422S1	CTTCTGCAAT CTTTTCGGTAA GCTCCCAAA CAATGGCCCTG GAGGCTTACG TCTTATGATCT CTTTTCGGTAA GCTC ----AG CAATGG--GG GAGGGGTGGG	1944 781
Pig Oct-4 promoter Mus S58422S1	TCTCTTATAGAG GGGGACGG AATTCCTT--A GG--TCCCAG TTGGGCGGC TTGATCAGCA TCAACACCGG AATTCCTTGA GGAAGGAG CAGGGTATCT	1988 830
Pig Oct-4 promoter Mus S58422S1	CTCTCTGAGG CTCGTCTTTT GAGGGAATGG GGGTGTGTTT GTTGGAGCTC CTCTCTGAGG CTCGTCTTTT GAGGGA-- --GG-----TGGAG---	2038 864
Pig Oct-4 promoter Mus S58422S1	TTTATGCTGC TCTGAAGGGG ATTTGTGTGTG AGGGGATGG GGCT--CTG ----AGCTG--GGGAAG-- --TTTGTGTG AGGGGATGG GGCTCAGCTG	2084 904
Pig Oct-4 promoter Mus S58422S1	GGGGTTGGGG TGCAGGAAGC TGTCCCCCAGG GGAGCCATCC TGGCCCATTC GGGGTTGGGG TGCAGGAAGT TGTCCCCCAGG GGAGCCATCC TGGCCCATTC	2134 954

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FIG. 10C

Pig Oct-4 promoter Mus S58422S1	AAGGGTTGAG AAGGGTTGAG	CACTTGTTTA GGGTTAGAGC TGCCCC CACTTGTTTA GGGTTAGAGC TGCCCC	TCT GGGGACCAGG TCT GGGGACCAGG	2184 1003
Pig Oct-4 promoter Mus S58422S1	ATTGTCCAGC CAAGGCCATT GTCC ATTGTCCAGC CAAGGCCATT GTCC	GCCCC CTTCGCCCCAG GCCCC CTTCGCCCCAG	TCCCTCCCA TCCCTCCCA	2234 1053
Pig Oct-4 promoter Mus S58422S1	GCCCTTTGA ACCTGAAGTC AGATATT GCCCTTTGA ACCTGAAGTC AGATATT	TCTCTCTCC TCTCTCTCC	CTCTCCCT CTCTCCCT	2282 1103
Pig Oct-4 promoter Mus S58422S1	CCCTTGGTTT TCCACCCCA GGGCTAGGG CCCTTGGTTT TCCACCCCA GGGCTAGGG	GTGGAAGCC GTGGAAGCC	AGATTGGAG AGATTGGAG	2331 1152
Pig Oct-4 promoter Mus S58422S1	GTGGGGGAGG GAGAAC GTGGGGGAGG GAGAAC	CTAGATATT CTAGATATT	GGGTCCCTGA GGGTCCCTGA	2381 1168

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INTERNATIONAL SEARCH REPORT

In: International Application No

PCT/US 98/21289

A. CLASSIFICATION OF SUBJECT MATTER

IPC 6 C12N15/00 A01K67/027 C12N5/06 C12N5/10 C07K14/47

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 6 A01K C12N C07K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

C. DOCUMENTS CONSIDERED TO BE RELEVANT

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Further documents are listed in the continuation of box C.



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Date of the actual completion of the international search

24 February 1999

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INTERNATIONAL SEARCH REPORT

In: International Application No

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C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

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In tional Application No

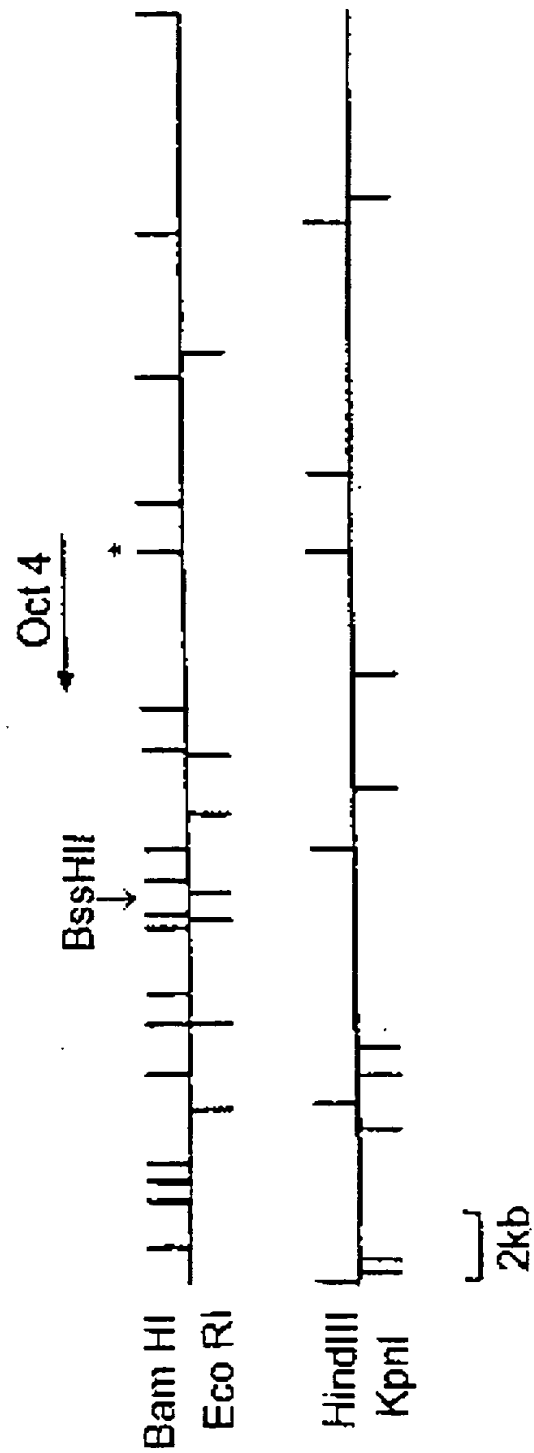
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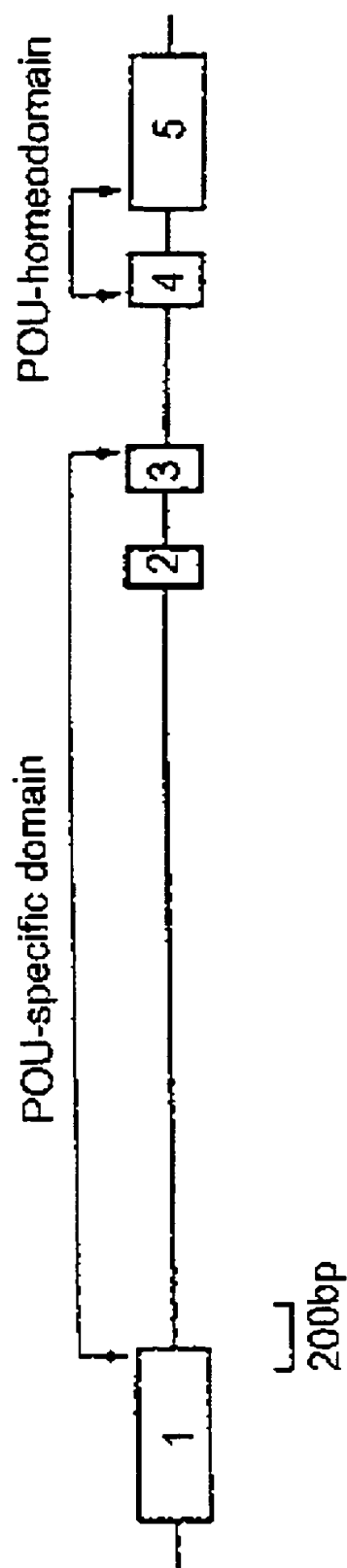
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FIG. 1A




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FIG. 1B



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FIG. 2A

BamHI SmaI PstI EcoRI 


AGTGGATCCCCCGGCTGCAGGAATTCGGG

BamHI

1 ATGGATCCTCGAAGCTGGCTAGCTTCCAGGGCTCCAGGTGGGCTGGATCGGACCA 60
 M D P R T W L S F Q G P P G G P G I G P -


21 CGCTCAGAGGTATTGGGGATCTCCCCATGTCCGCCCATACGAGTTCCTCGGAGGGAIG 120
 G S E V L G I S P C P P A Y E F C G G M -

41 GCATACTGTGGACCTCAGGTTGGTCTGGGCTAGTCCCCCAAGTTGGCGTGGAGACTTG 180
 A Y C G P Q V G L G L V P Q V G V E T L -

 Clone 16

61 CAGCCTGAGGGCCAGGAGCAGGAGTGGAAAGCAACTCAGAGGGAACCTCTCTGAG 240
 Q P E G Q A G A R V E S N S E G T S S E -

81 CCTGTCCGAGCCGCCATGCCGTGAAGTTGGAGAGGTGGAAACCACTCCCGAGGAG 300
 P C A D R P N A V K L E K V E P T P E E -

PstI  Clone 1

101 TCCGAGACATGAAGCCCTGCAGAGGAGCTAGAACACTTTCCTGCTGAGGAG 360
 S Q D H K A L Q K E L E Q F A K L L K Q -

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FIG. 2B

POU-SPECIFIC DOMAIN

121 AAGAGGATCACCTTGGGGTACACCCAGCGGACGTGGGGCTCACCCCTGGGGCTTCTCTT 420
K R I T L G Y T Q A D V G L T L G V L F -

141 GGAAGGTGTTCAAGCCAGACCCATCTGTGGCTTCGAGGCTTGACGCTCAGCCTTAAG 480
G K V F S Q T T I C R F E A L Q L E L K -

161 AACATGTGTAAAGCTGGGGCCCTGCTGGAGAGTGGGTGGAGGAGCCGACACAAATGAG 540
N H C K L R P L L E K W V E E A D N N E -

181 AACCTTCAGGAGATATGCAATCCGAGACCCCTGGTGCAGGCCCGGAGAGAACCGAACT 600
H L Q E I C K S E T L V Q A R K R K T -

201 ACCATTGAGAACCGTGTGAGGTGGAGTCTGGAGACCAATGTTCTGAAGTGCCCGAGCCCC 660
S I E H R V R W S L E T H F L K C P K P -

221 TCCCTACAGCAGATCACTCACATCGCCCATCGCTTGGCTAGAGAGATGTGGTTCGA 720
S L Q Q I T H I A N Q L G L E K D V V R -

241 GTATCGTTCGTAAACCGCGCCAGAGGGCAAGATCAAGTATTGAGTATTCGCCACCA 780
V W F C N R R Q K G K R S S I E Y S Q R -

POU HOMEODOMAIN

FIG. 2C

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GAAGAGTATGAGGCTACAGGACACACCTTCCAGGGGGGCTGTATCCCTTCCCTCTGCC 840
 261 E E Y E A T G T P F P G G A V S F P L P -

 CCAGGTCCCCACCTTTGGCACCCCAGGCTATGGAAGCCCCACCTTCACACACCTACTCA 900
 281 P G R H F G T P G Y G S P H F T T L Y S -

 GTCCCTTTCCCTGAGGGGAGGCCCTTCCCTCTGTCCCGTCACTGCTCTGGGCTCTCCC 960
 301 V P F P E G E A F P S V P V T A L G S P -

 ATGCATTCAAACTGAGGCACCCAGCCCTCCCTGGGATGCTGTGAGCCAGGCAAGCGAGG 1020
 321 H H S N * 324

 TAGACAAAGAGAACCTGGAGCCTTTGGGGTTAAATTCTTTACTGAGGAGGGATTAAAGCA 1080
 CAACAGGGGTGGGGGTGGGATGGGGAAGAAGCTCAGTGATGCTGTGATCAGGAGCCT 1140
 GGCCTGTCTGTCACTCATCATTTTGTGTTCTAATAATAGCAGTGGACACACAGTAAAAA 1200
 AAAAAAATACTCGAG 1217
 XhoI

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FIG. 3A

10	20	30	40	50
* * *	* *	* *	* *	* *
GAATTCGGCT TCACCAGGCT TCGGACTTCG CCTTCTCGCC CCGCCGGGC				
60	70	80	90	100
* * *	* *	* *	* *	* *
GGTGGAGCG ATGGGCCGGG AGGGCGGAGC CGGCTGGGT TGATCCTCGG				
110	120	130	140	150
* * *	* *	* *	* *	* *
ACCTGGCTGA GCTTCCAAGG GCCTCCCGGT GGGTCAGGA TCGGCCCGG				
160	170	180	190	200
* * *	* *	* *	* *	* *
GGTTGGCCG GCGGCCGAGG TGTGGGGCT TCCCGGTGC CCCCCCCT				

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FIG. 3B

210	220	230	240	250
* * *	* * *	* * *	* * *	* * *
ATGACTTCTG	CGGAGGGATG	GCCTACTGCG	CACCTCAGGT	CGGAGTGGGG
260	270	280	290	300
* * *	* * *	* * *	* * *	* * *
CTCGTGCCCC	AGGGCGGCCT	GGAGACCCCT	CAGCCCCGAGG	GCGAGGCGGG
310	320	330	340	
* * *	* * *	* * *	* * *	
GGCCGGGGTG	GAGAGCAACT	CCGAGGGGGC	CTCCCCCGAG	

FIG. 4A

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10	20	30	40	50
* * *	* * *	* * *	* * *	* * *
GGATCCTCGG	ACCTGGCTGA	GCTTCCAAGG	GCCTCCCGGT	GGGTCAGGGA
60	70	80	90	100
* * *	* * *	* * *	* * *	* * *
TCCGGCCCGG	GGTTGGCCCG	GGCGCCGAGG	TGTGGGGGCT	TCCCGCGTGC
110	120	130	140	150
* * *	* * *	* * *	* * *	* * *
CCCCCGCCCT	ATGACTTCTG	CGGAGGGATG	GCCTACTGCG	CACCTCAGGT
160	170	180	190	200
* * *	* * *	* * *	* * *	* * *
CGGAGTGGGG	CTGGTGCCCC	AGGGCGGCCT	GGAGACCCCT	CAGCCCCGAGG

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FIG. 4B

210 * * * * * 220 * * * * * 230 * * * * * 240 * * * * * 250 * * * * *

GGAGGGGG GGCCCGGTG GAGAGCAACT CCGAGGGGC CTCCCCCGAG

260 * * * * * 270 * * * * * 280 * * * * * 290 * * * * * 300 * * * * *

CCTGTGCCG CCCCCGCTGG CGCCGCGAAG CTGGACAAGG AGAAGCTGGA

310 * * * * * 320 * * * * * 330 * * * * * 340 * * * * * 350 * * * * *

GCCGAACCC GAAGAGCGA GTGAGCTGCC GGGAGCTGGG GGAGGCGATC

360 * * * * * 370 * * * * * 380 * * * * * 390 * * * * * 400 * * * * *

GCGTGCCG GGGCGCACG CAGGGAGGT GTCCGCTGC CCCCCGGCA

*

GGAGGGG

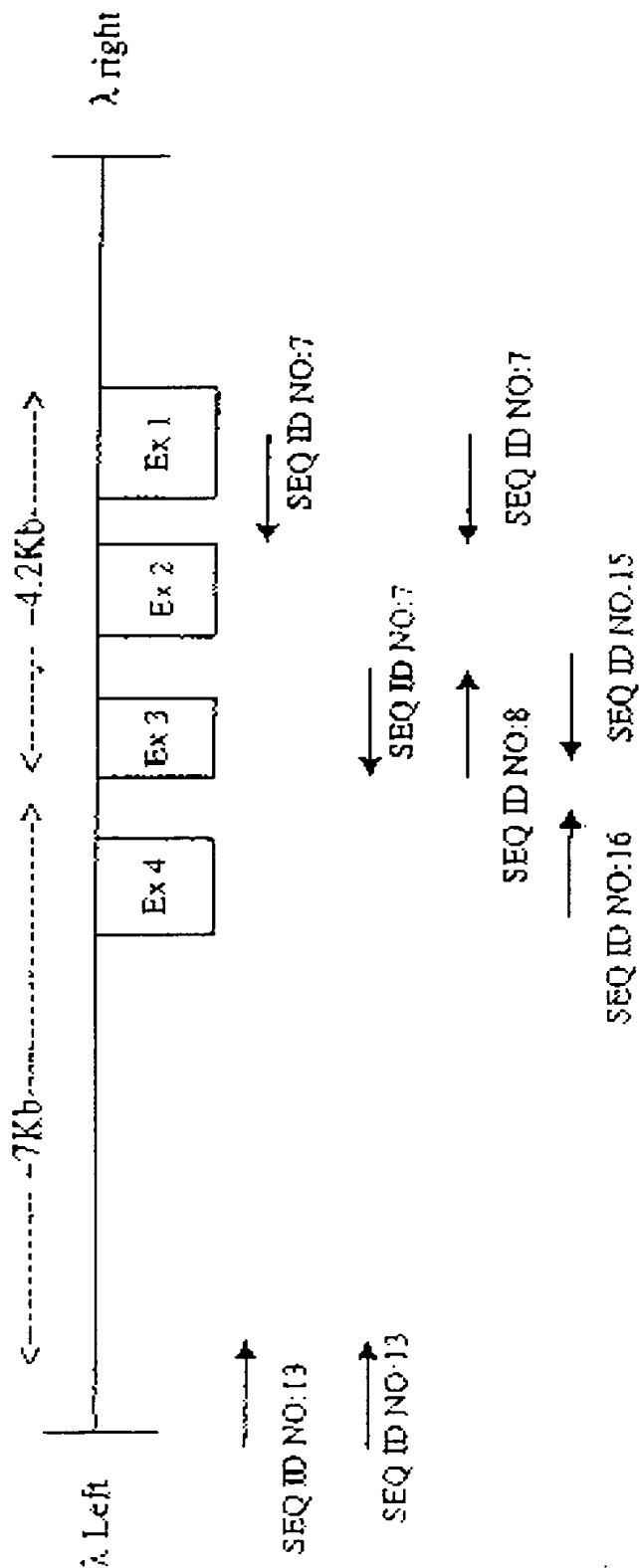
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FIG. 5A

Oct-4 CLONE 3



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FIG. 5B

Oct-4 CLONE 4

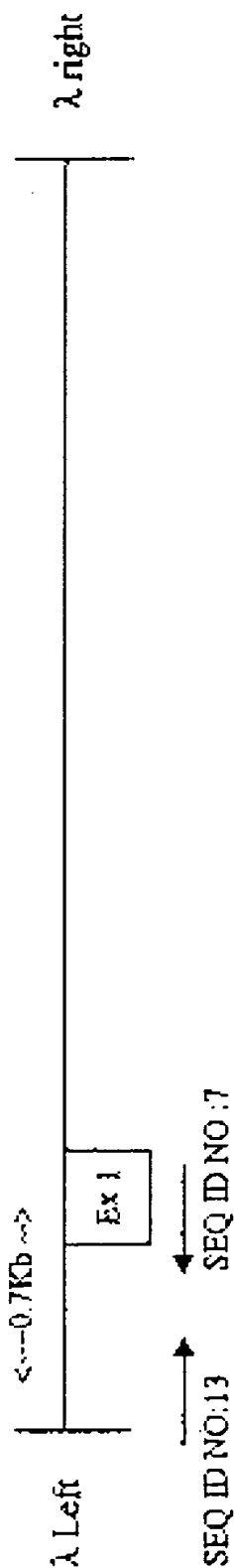


FIG. 6A

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10	20	30	40	50
* * *	* *	* *	* *	* *
AGCCATGCTG	GGTTGATCCT	CCCACCTGGC	TGTGCTTCCA	AGGGCCTCCT
60	70	80	90	100
* * *	* *	* *	* *	* *
GTGGGTCAG	GGATCTGGCG	GGGGGCTGCT	GGACCCAGAG	GTGGGGAGGC
110	120	130	140	150
* * *	* *	* *	* *	* *
TTCTCTCATG	CCCCCGGCC	TAGGACTTCT	GCGGAGGGAT	GGCCTACTGT
160	170	180	190	200
* * *	* *	* *	* *	* *
GCACCTCAGG	TCAGAGAGGG	GCTGGTGCCC	CAAGCGGGCC	TGGAGACCCC

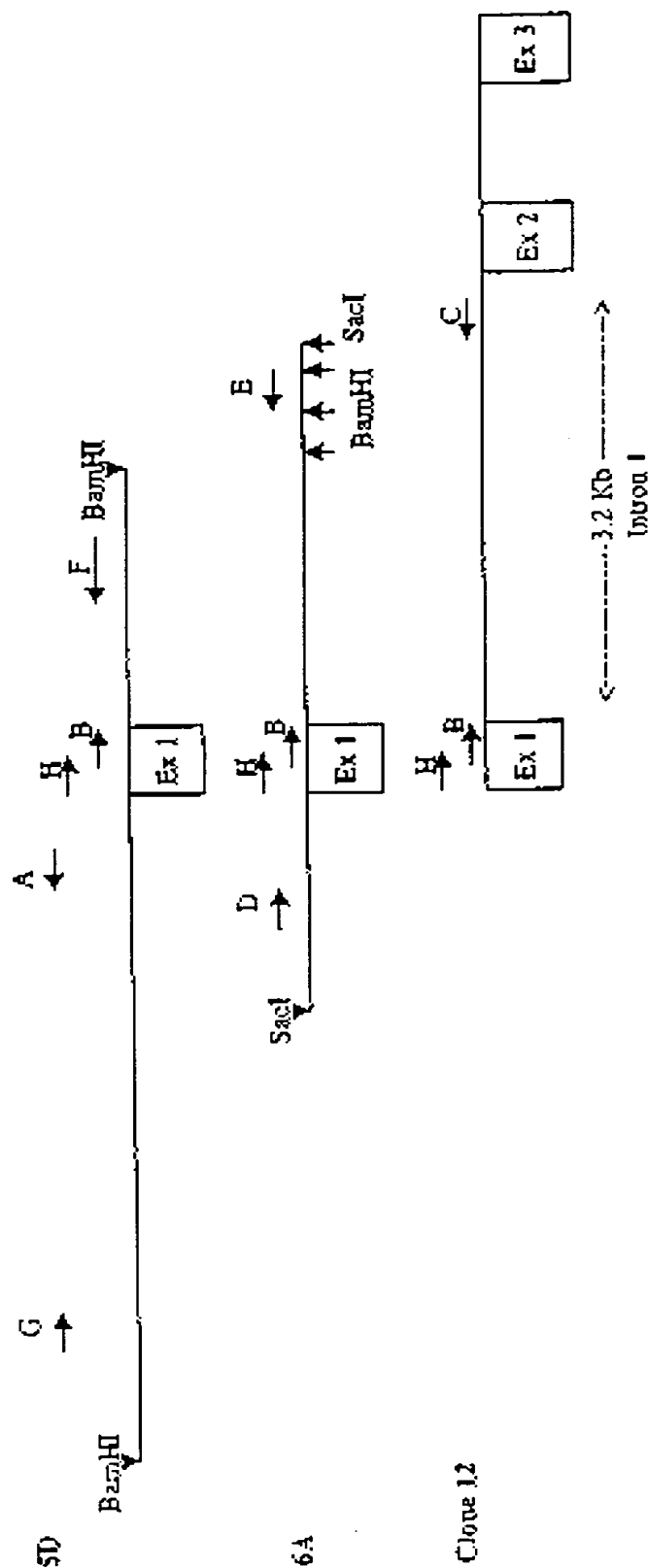
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FIG. 6B

210	220	230	240	250
* * *	* *	*	*	*
TCAGCCCTAG	GGCCAGGCAG	GAGTCGGGGT	GGGAGCAAC	TCCGAGGGGG
260	270	280	290	300
* * *	* *	*	*	*
CCTCCCTGGA	GCCCTATGCC	ACCCCGGTG	GCACTGCACA	GCTGGACAAG
310	320	330	340	350
* * *	* *	*	*	*
GAGAAACTAG	AGCCGAATCC	TGAGAAGTCC	CAGGACATCA	AAACGCTTCA
360	370			
* * *	* *	*		
GAAAGACCTT	CAACAAATTG	CCAAGCTT		

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FIG. 7



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FIG. 8A

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GGATCCCTAG CCTGGAACCC TCCATAAGCC GTGGTACTG CCCTAAGAAG AAAAAAAG 60
TGGTTTCTTA CCTTGGTCTA GAGCAAGCCT CCATTTTCCC CAGGAGTCAT TTCAGCTGGT 120
TTTCCCTACC AAACAGCAAG GGGATGGCCG GGCTGACAGC AGCAAAGTCA CTGTCACTC 180
TTTGCAGGCT TGCCAGGCCA GCTGCATCTG GCAGGGAGCG GCAGCTCTCA CCTGCCCTCC 240
CTGGGTCATG CTCTACAGCC AGATACTTGG CATTTGTCTT TGTGTAGGCG CTCAATATTG 300
TACTCTAATA AGGGTACATG TGGGAGTTCC CTGGTGGCTC AGCAGTTGAG GATCTGGCAT 360
TGTCRCTGCT CTGGCATGGA TCTCTGCTCT GCGCGAGGTT CAATTCCCTGG CCTGGGAACT 420
TCTGTACGCC GCAGGCGTGA CTGGAAAAAT ACAGGTGGGG TGGGGTGAGG AGTGTATGTG 480
GAGAGTCTGC AAACCCAGGC CTAAATTGGT TTGGGGGACT TGAAGTTTAT AGTGACTCCC 540
TACCCMAAAG AGTGGAGMAG CCAGGTCTGA TGACTTAAAC CCACCTGCAG TCTGCTCTGG 600
GCCTGCAGAG ACCTGGCCTC TGCAAGAGTG AAGCTGCCTA CACTTCAGGC CTACACGGAG 660
GGTGGGGAGG AGAGGGGAAT AGGCTCAGCC CTGCCATGCC AAGCACCCCC AGGCTGACTA 720
GGACTCCAGA CAAATTTAGC TTGTCTCTTAA GGTTCCTGGT CAGACCCCCAG GCAAGCACAG 780
AACTGATCTG GCTCAGATGT CTGGCTACAA GCTATCCAGG AACCACGGCA TCCAGCCCTC 840
CCCAGCCCTC CCCAGGCTTT CCTCTTGGA TAGGAAGGAC ACTTGTCTTAA ACCAGAAACA 900
TACCATCTAG AGCAGCTATT TATGGTGTAT TAAAAACAC AGGGTGCTAT TTAGTCGGGG 960
GTGGGTGGGA AGGAGAGAGG TGTTTAGGGT CCGCGGGGAA GTCAGGGGCA CAGGGGCTCT 1020
CTGGACCCACA TGGGGAGAGG GGTTCCTGGG AGGCCAGAGG GCGAGGAGCC AAGGAGCTCA 1080

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FIG. 8B

GCAGTAGATT CCTAGGCCC GCCCTTCCCC CTCCTCAGGG AGGCCGTCTT CTYGGCAGAC 1140
 AGCAGATAGA TGCATGACAA AGGGCCCATG ATGGCTCTGT CCTGGGGGTT GGGGAGATGG 1200
 CTAGGGAGGG GCCCCTCCTG GTCTGAAGCA CATCTTTCCA CCCCACCAG GCCCCTTAAT 1260
 CTATCTGCTT TTGGGGCAGT TAGTAGTTTA GAGTTGAAT AGCTCCAGCC CTGCTGCCCT 1320
 ATAAATCTTT CAACAGACCT ATGGGAAGTA TTGAATGCA TGCACGCAAT TAGTCACCCC 1380
 AAACGCACAG GCCGATGGC ACTGGAAGAG ATTCAGAGGA GAAAAGCAA AACAAACAA 1440
 CACGGACACA CAAAACCCA ACAGACTCAA AGGACTCCTG GTGGAGCTAA CTGGTCACAG 1500
 TCTGGAGGAT GCCAGCCCCT CAAGACAGAT GCCGAGCCAC TGACCCCTAGC AAACAACCTC 1560
 AGACCCAGCC AAGATGAAGA GGTGTCCTAGG TCCGCAGAGG TCTGTGTCCC AGTCTCAGGA 1620
 GTCTGGCCTC CAACTGTAG GAAGCTCTGA TCCATGGCTT CTCIGGAGAG CCCCCCTCAC 1680
 TCAGGTTTAC CTGGGGCCTT CGTTTAGGGC AAGTTGGGG AGCAGACAGA CAAACATCAT 1740
 CCCCAGCAGA CAGCCAGTCT GAAGCTATT CTCGTGCAA CAGATCAG CACTAGGCCA 1800
 GCAGCCTGAG CCTCAGGACA GACCCAGAAA AATAGACCCT GTGGGAGAGC TTAGGGCAGG 1860
 ATTCTGTGAC CCCCCTCCCC ATCGCAGTTC ACCCCCTTCT GCATCTTTTC GCTAGCCCCC 1920
 CAAACAAAGG CCTGGACGCC TCAGTCCCTCT AGAGGGGGGA CAGGATACCT AGGTCCCAGT 1980
 GGGGGGCCCT GTCTGAGGCT CAGTCTTTGA GGGGATGGGG GTCTTGTTC TGGAGCTCTT 2040
 TTAGCTGCTC TGAAGGGAT TCTGTGTGAG GGGATTGGGG CTGGGGGTT GGGGGCAGG 2100
 AAGCTGTCCC CAGGGGAGCC ATCCAGGCC ATTCAAGGGT TGAGCACTTG TTIAGGGTTA 2160

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FIG. 8C

GAGCTGCCCC CTCGGGGAC CAGGATTGTC CAGCCAAGGC CATGTGTCGG CCCCCTTCCC 2220
 CCAAGTCCCTC CCAAGCCTCT TTGAACCTGA AGTCAGATAT TTTTCTCTCC CCCCCCTCC 2280
 CTCCTTGGCT TTCCCCAACC AGGCCCTAGG GGTGGAGGCC CAGATTGGGA GGTGGGGAG 2340
 GGAGACAGT CAACTATGG GCTAGATATT TGGTCCCTG AAGGGGGCT GGGGACAAAG 2400
 GAACCTGATG TCGCGCGGGA CCCACAGCG GGGACCTGCG AGCGGGTGTG CGATTGATT 2460
 CTCGTCCCTGC ACRAAGATTG GGAGACTCAG GCCCAGTCCA TOGAGCTTGA TCCCTGGGAG 2520
 GGAATAATGG GGTTCATCC CTCGGTCTGG TGGAAAGGAG GCCCCGGMAC CCGGAATACT 2580
 GTACGGAAATG GAGCCCCGTG TGGCAGTCNG CCCCTGGTGA GGGGTGGAAT CTAATAGGCT 2640
 GGGCGGATGG TTGCTGGGA TCCAGCTTT GGGGTGCCGG AATCTGGCCA GTAATCTAGT 2700
 TGGGAATGCC TAGGTTCCCG GACTGGCGT GAGGSCAGAG AGCAGGAAT GAGGAGTAGC 2760
 TCCGGCAGGA CTTAGCACAG ACACCAAGCC TGTGTGAGGA CCTGAGAGGG TCGCTGGGCT 2820
 CCCTTGAGGA GACAGTCCCA GGTCTCTGA AGAGGGGTCC AACACCTGGC TCCCAGACAG 2880
 CCCCRAATGT CACAGAGCAG TGGAGAGGGC CGGGCGGCGG GTTGGAGTT GGAGGTGAG 2940
 GCGGCATGGG GGAACCTGAC CAGGGGCTG GGGACCCGAG AGCGGCCCGG GCGGACCTCT 3000
 CCGACTTTGG CCGTCCAGAC ACCACCGCCA CCAGCCAGCA AACACCTCC GCTCAGTTT 3060
 CTCCCACCCC CACCGACCC TCCCCAACC ATCCAGGGG CGGGGCCAGA GGTCAAGGT 3120
 AGTGGGTGG ATTGGGGAG GAGAGAGGTG TOGAGCAGTC CCTTGGAGA GCCCTGCTT 3180
 TACTGGGCC CCGGCTTGG GGGCTTCTT TCCCC 3215

[illegible]

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FIG. 9B

pig oct4 2701-3215	C-AGAGAGCTT	CGAAGAGGG	TCCATCA--C	CTGCTCTCC	EACAGCC	185
huOct4 1-499	T-CAGAGCTT	CAATCAG--	CCATCP--T	TGTACTCCPC	TGCA-CTCA	181
mOct4 1401-1950	CCAGAGCTT	C---AGAG-	CCATCPGG	TTTGTCTTC	EATGCGG--	225
pig oct4 2701-3215	ATGTGACAT	AGCAGTGGG	AGG-GCTTGG	CGCCCGGTG	GGAATTGAG	234
huOct4 1-499	CTGTGCTCA	CAAGCTTGG	A---CCCTT	C---TTAATAA	ATTAATAAA	225
mOct4 1401-1950	ATGTGACAT	CTGCTTGGG	AGGTGCTTGG	TGACCTCA-AG	GCTGGGGTGA	274
pig oct4 2701-3215	GTGATGGCG	CATGCTTAC	CTGATCCAG	GCTCTGCTGA	CCGAGAGGG	284
huOct4 1-499	AAATGCTTC	TGTGCTTAC	CTGATCTAG	GCTCTGG--	---AGGGG	268
mOct4 1401-1950	GACATGCTT	AGCTTCTTA	ATGATGGCC	TTCCTGCGT	CC-----C	316
pig oct4 2701-3215	GAAGGCTGG	ACCTTCTTA	CTTCTGCTT	C---CAAGAC	CACTTACC	332
huOct4 1-499	GTCAGTCTT	GTCTTCTT	TTTCTCTCT	CTAATCTAC	CACTTACC	316
mOct4 1401-1950	GACTTATGG	GTATGCTT	TTCTGCTT	CTCTA---AC	CTCTCTTGG	360
pig oct4 2701-3215	ATC---CAT	CAATCTCT	TCTGCTCTAG	TTTCTCTAC	CCCTCTGAC	377
huOct4 1-499	ATCAT-TAG	CAATCTCT	TTTCTCTAG	TTTCTCT--	CTCTCTTCC	362
mOct4 1401-1950	ATGACATCT	CAATCTCT	CTGCTCTAG	TTTCTCTACC	CCCTCTTCC	408

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FIG. 9C

A

pig oct4 2701-3215	CT-CTCCCTCC ACCCAATCCAG GGGGGGGGGC CAGAGGTCAA GGCTAGGCGG	426
huOct4 1-499	CT-CTCCCTCC ACCCAATCCAG GGGGGGGGGC CAGAGGTCAA GGCTAGGCGG	411
moOct4 1401-1950	CTCTCCCTCC ACCCAATCCAG GGGGGGGGGC CAGAGGTCAA GGCTAGGCGG	458
pig oct4 2701-3215	TGGGATGGG GAGGGAGAGA GGTGTGATCC ATCCCTTTTG GAGAGCCCTG	476
huOct4 1-499	TGGGATGGG GAGGGAGAGA GGTGTGATCC ATCCCTTTTG CA-AGCCCTTC	460
moOct4 1401-1950	TGGGATGGG GAGGGAGAG- ---GTATATC GTCCCTTTTG ATCAGCCCTTC	503
pig oct4 2701-3215	CTTTTACTTG GCGCGCGGCT TGGGTTCCTT TCCCTTCCCG	515
huOct4 1-499	ATTTCACCTTG GCGCGCGGCT TGGGTTCCTT TCCCTTCCCG	499
moOct4 1401-1950	TTTTCACCTTG GCGCGCGGCT TGGGTTCCTT ACCCTTCCCG TGGCTGG	550

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FIG. 10A

Pig Oct-4 promoter Mus S58422S1	<p> TCACAGTCTG GAGGATGCCA GGGCTCAAG ACAGATGCG AGCCACTGAC ----- 3 AGCCACTGAC </p>	1544 440
Pig Oct-4 promoter Mus S58422S1	<p> CCTAGCAAC AACTCAGAC CCGCCAACTA TGAAGAGG TGAAGAGG CCTAGCAAC AACTCAGAC -- -- -- -- -- -- -- -- -- -- -- -- -- </p>	1594 475
Pig Oct-4 promoter Mus S58422S1	<p> CAGAGTCTG TGTCCGCTC TTAGGAGTCT GGGCTCAAA CTGAGGAG CAGAGTCTG T -- -- </p>	1644 510
Pig Oct-4 promoter Mus S58422S1	<p> CTCCTATCCA TGGCTTCTT GAGGAGTCT CATTATCAG GTTCCTCTG CTCCTG -- -- </p>	1694 545
Pig Oct-4 promoter Mus S58422S1	<p> GGGCTTGGTT TAGGGCAAT TGGGAGCA GACGACAAA CATTATCCC GGGCTTGGTT TAGGGCAAT TGGGAGCA GACGACAAA CATTATCCC </p>	1744 595
Pig Oct-4 promoter Mus S58422S1	<p> TSCAGACAGC CATTCTGAAA CCTATCTCT TGCATAAATA ATCAGCACCT TSCAGACAGC CATTCTGAGG CCTATCTCT TGCATAAATA ATCAGCACCT </p>	1794 645

FIG. 10B

AGGCGCAGCA	TCTTTC	AGACTGGCC	CAGAAATATA	EACCTTTTG
AGGCGCAGCA	TGTGTTC	AGACTGGCC	CAGAAATACA	-----CTTTTG

CTGCACCCC TCCACATCC CATTTCACCC
CTGCACCCC TCCACATCC TCCATTCCTGTC

CGTTCTGAT CTTC CGCCTA GC CCCCATA CAAAGCCCTG GAAGCTTAA
TATTACACT CTTC CGCCTA GGC-----AG CAATGG--GG GAAGCCGTGG

TCTCTTAGAG GCGGACGGG A TTTCTT--A GG--TCCGAG TCGGGGGC
 TTGGACCTAG TCAACACGGG A TTTCTTGA GCGAGGGAG CCGGGATATT

CTCTCTGAGG CTCTGCTTT GAGCGAGGG GCGGTGTTT GGTGGAGTTC
 CTTCTGAGG CTCTGCTTT GAGCGAGG GCGGTGTTT GGTGGAGTTC

TTTTAGCTGC TCCTGAGGG PTTCTGTGTG AGGGGATTGG GCGT --- 3-6
 --- AGCTG --- GGTGAGC --- TTTCTGTGTG AGGGGATTGG GCGT TAGGTC 3-6

52GGTTGGGG	53CAGGAAGC	TGTCCCAGG	GGAGGCATCC	52GGCCATTCC
53GGTTGGGG	53CAGGAAGC	TGTCCCAGG	GGAGGCATCC	52GGCCATTCC

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FIG. 10C

Pig Oct-4 promoter Mus S58422S1	AAGGGTGGAG AAGGGTGGAG	CACTGTGTTA GGGTTAGAGC TGCCCC CACTGTGTTA GGGTTAGAGC TGCCCC	TCT GGGGACCAGG TCT GGGGACCAGG	2184 1003
Pig Oct-4 promoter Mus S58422S1	ATTGTCCAGC CAAGGCCATT GTCCGGCCCC ATTGTCCAGC CAAGGCCATT GTCCGGCCCC	CTTCCCCCAG TCCCTCCCA CTTCCCCCAG TCCCTCCCA		2234 1053
Pig Oct-4 promoter Mus S58422S1	CCCTCTTTGA ACCTGAAGTC AGATATT CCCTCTTTGA ACCTGAAGTC AGATATT	T TCTCTCTCTCC T TCTCTCTCTCC	CTCTCCCT-T CTCTCCCTTC	2282 1103
Pig Oct-4 promoter Mus S58422S1	CCCTGTGTTT TCCACCCA GGGCTAGG CCCTGTGTTT TCCACCCA GGGCTAGG	GTGGTGGCC GTGGTGGCC	AGATTGGGAG AGATTGGGAG	2331 1152
Pig Oct-4 promoter Mus S58422S1	GTGGGGGAGG GAGAACGTC GTGGGGGAGG GAGAAC	AACTATGGGG CTAGATATT -----	GGGTCCCTGA -----	2381 1168

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